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14:40:13 2004
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isogene, which may be done by turning off by transforming a targeted organ, tissue or cell population with an expression vector that expresses high levels of untranslatable mRNA for the isogene. Specific therapeutics identified by these methods may be useful for allergic diseases. The present sequence is a probe for human ILAR-alpha 88888888

Sequence 15 BP; 5 A; 4 C; 5 G; 1 T; 0 U; 0 Other;

Gaps .; Length 15; Query Match
13.4%; Score 9.8; DB 1; Length 15
Best Local Similarity 84.6%; Pred. No. 1.2e+03;
Matches 11; Conservative 0; Mismatches 2; Indels 900 CCTGGTCATTTTC 912 à d

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15

RESULT 1332 AAS98327

AAS98327 standard; DNA; 15 AAS98327;

BP

(first entry) 12-MAR-2002

Galanin receptor gene GALR1 allele-specific oligonucleotide #39.

SNP; Galanin receptor; GALR1; human; single nucleotide polymorphism; drug discovery; haplotyping; infectious diarrhoea; growth hormone deficiency; allele-specific oligonucleotide; ss.

Homo sapiens

WO200179237-A2.

25-OCT-2001.

16-APR-2001; 2001WO-US012306

14-APR-2000; 2000US-0197838P.

(GENA-) GENAISSANCE PHARM INC.

봈 Denton RR, Nandabalan Choi JY, Bentivegna SC, Chew A,

WPI; 2002-066341/09.

Genotyping human galanin receptor gene of an individual for determining haplotype of an individual, involves determining the identity of nucleotide pair at specific polymorphic sites for two copies of the gene.

Claim 16; Page 15; 99pp; English

The invention relates to genotyping human galanin receptor (GALRI) gene

of an individual, involving determining for the two copies of the GALRI

come present in the individual, the identity of the nucleotide pair at

one or more polymorphic sites. The method is useful for determining

whether an individual has a haplotype or haplotype pairs defined in the

specification. This is useful for improving the efficacy and reliability

of several steps in the discovery and development of drugs for treating

diseases associated with GALRI activity, e.g., infectious diarrhoea and

creating a specific condition or disease predicted to be associated with

CALRI activity, and in the design of clinical trials of candidate drugs

for treating a specific condition or disease predicted to be associated

with GALRI activity. The method is useful to screen for compounds

creating GALRI to treat a specific conditions or disease associated with

GALRI activity. A GALRI polymucleotide or variant is useful in studying

the expression and function of GALRI, and in expressing GALRI protein for

use in screening for candidate drugs to treat diseases related to GALRI

cativity. The polymucleotide or variant is useful for studying expression

of the GALRI isogenes in vivo, for in vivo screening and testing of drugs

targeted against GALRI protein, and for studying the effect of the

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variation on the bological activity of GALR1 as well as on the binding affinity of candidate drugs targeting GALR1 for the treatment of infectious diarrhoea and growth hormone insufficiency. AAS98289- AAS98408 represent human GALR1 gene allele-specific oligonucleotides used to detect GALR1 gene polymorphisms as described in the method of the invention
                                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                           Query Match 13.4%; Score 9.8; DB 1; Length 15; Best Local Similarity 73.3%; Pred. No. 1.2e+03; Matches 11; Conservative 1; Mismatches 3; Indels
                                                                                                                                                                                   G; 3 T; 0 U; 1 Other;
                                                                                                                                                                                   Sequence 15 BP; 1 A; 7 C; 3
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RESULT 1333 ABK97325,

ABK97325 standard; DNA; 15 BP.

ABK97325;

07-OCT-2002 (first entry)

16S rRNA gene B-C synthetic variant PCR primer.

Strain identification method, prokaryote, eukaryote, ribosomal DNA; HCR, highly conserved region, highly variable region; HVR; bacterium; methicillin-resistant Staphylococcus aurents; nosocomial infection; ss; DNA fingerprinting; pathogenic bacteria; infection control; PCR; primer; restriction fragment length polymorphism; RFLP; 16s rRNA; 23s rRNA; 5S.

Synthetic.

JS6395475-B1

28-MAY-2002,

95US-00461210. 05-JUN-1995; 93US-00064596.

18-MAY-1993;

(UYFL) UNIV FLORIDA STATE

Reeves Whitehouse E, Leggett CG,

RH;

WPI; 2002-556092/59.

Identifying strain of prokaryote or individual of eukaryote, useful in clinical laboratories for strain identification of pathogenic bacteria, comprises amplifying specific DNA fragment in ribosomal RNA intergene region.

Disclosure; Col 5; 31pp; English.

The present invention relates to a new method of identifying strain of prokaryote or individual of eukaryote. This method involves amplifying a highly conserved region (HCR) of ribosomal DNA of prokaryote or eukaryote, where the HCR of DNA flanks a highly variable region (HVR) of DNA, to generate amplified DNA fragments that are separated fragmented to yield labelled, amplified DNA fragments that are separated by electrophoresis so that prokaryote or eukaryote can be identified. The invention can be used for identifying a strain of a prokaryote or an individual of an eukaryote. The method is proferably useful for identifying a prokaryote. The method is proferably methicillin-resistant Staphylococcus aureus. The method is useful for identifying different bacterial strains involved in e.g. nosocomial infections, and for identifying species such as pedigrees, with respect to a enkaryote. The method is sensitive enough to detect cespect to a enkaryote. The method is sensitive enough to detect differences between e.g. bacterial isolates of the same species. The

Sequence 15 BP; 5 A; 3 C; 6 G; 0 T; 0 U; 1 Other;

gene polymorphisms

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methods generally depend upon rapid, semi automated DNA analysis, and more particularly, upon a type of DNA fingerprinting of multiple segments of DNA. The methods are beneficial in clinical laboratories, because they allow for rapid strain identification of pathogenic bacteria. The method is more definitive since genomic bacterial DNA is used. The method also provides results with great speed e.g. a preliminary screen by agarcsee gel electrophoresis of a polymerase chain reaction (PCR) product can be completed 5-6 hours after receiving hospital isolates. The preliminary screen can then be confirmed in approximately 24 hours by restriction fragment length polymorphism analysis (RPLP). The speed of the methods provide infection control personnel with adequate information to contain analysis done retrospectively. The present nucleic acid sequence represents one of a collection (RBS97222-ABK99726) of PCR primers used in the methods of the invention, as described above
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; single nucleotide polymorphism; SNP; IMPDH2; chromosome 3p21.2;
IMP dehydrogenase 2; haplotyping; genotyping; cancer; cytostatic;
allele-specific oligonucleotide; ASO; primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ASO primer #24 to detect IMPDH2 gene polymorphisms.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 15 BP; 6 A; 2 C; 5 G; 2 T; 0 U; 0 Other;
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AAS18277/C
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AC AAS18277/C
DT 25-FEB-:
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DT 25-FEB-:
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M HUMAN;
IN HORD;
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M HORD Sal
CC (SNP)
CC (SNP)
CC Almel-
CC Almel
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The present invention relates to genotyping protease inhibitor (PI) 4 (kallistatin) gene of an individual, involves determining for the two copies of the PI4 gene present in the individual, the identity of the nucleotide pair at one or more polymorphic sites. PI4 gene is located on chromosome 14931-q32.1. Genotyping is useful for determining if an chromosome 14931-q32.1. Genotyping is useful for determining if an specification. Haplotype or haplotype pairs defined in the specification. Haplotype or haplotype pairs defined in the challing of several steps in the discovery and development of drugs reliability of several steps in the discovery and development of drugs for treating diseases associated with PI4 activity, e.g. acute pancreatitis, to validate PI4 as a candidate agent for treating a specific condition or disease predicted to be associated with PI4 activity, and in the design of clinical trials of candidate drugs for treating a specific condition or disease predicted to be associated with PI4 activity. The PI4 gene is useful in studying the expression and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Genotyping protease inhibitor 4 gene of individual for determining haplotype of individual, involves determining identity of nucleotide pair at specific polymorphic sites for two copies of gene.
                                                                                                                                                                                                                                                                                                                                                                                                             Human, protease inhibitor, PI4; kallistatin; therapy; polymorphic site; PS; haplotyping; genotyping; acute pancreatitis; drug screening; antiinflammatory; chromosome 14q31-q32.1; primer; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  function of P14, and in expressing P14 protein for use in screening for andidate drugs to treat diseases related to P14 activity. The present sequence is a ASO (allale-specific oligonucleotide) primer to detect human P14 gene polymorphisms
                                               Gaps
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13.4%; Score 9.8; DB 1; Length 15; 73.3%; Pred. No. 1.2e+03; tive 1; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                      ASO primer #25 to detect human PI4 gene polymorphisms.
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                                                                                                                                                                                                                                         AAD25989 standard; DNA; 15 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               13-APR-2001; 2001WO-US012255.
                                                                                         912 CTITGGTCTTTGCCT 926
                                                                                                                             15 cygrecicicicii
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                                               11; Conservative
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Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Choi JY, Koshy B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO200179227-A2.
                                                                                                                                                                                                                                                                                                                             26-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens.
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                                             Matches
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Gaps

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13.4%; Score 9.8; DB 1; Length 15; 84.6%; Pred. No. 1.2e+03; ative 0; Mismatches 2; Indels

Query Match
Best Local Similarity 84.6
Matches 11; Conservative

The present invention relates to novel single nucleotide polymorphisms (SNPs) in the human IMP dehydrogenase 2 (IMPDH2) gene located on chromosome 3p21.2, and methods for haplotyping and/or genotyping the IMPDH2 gene in an individual. The methods of the invention make use of allele-specific oligonucleotides (ASOs) as probes and primers and/or primer-extension oligonucleotides for detecting the IMPDH2 gene polymorphisms. The polymucleotides and screened compounds are useful for (developing) treatment of diseases associated with IMPDH2 activity, such as cancer. AAS18254-AAS18279 represent ASO primers for detecting IMPDH2

Claim 15; Page 13; 70pp; English.

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942 CATTGGTTTAATG 954
             CATTAGATTAATG 1
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RESULT 1336

AAS98674 standard; DNA; 15

ВР

AAS98674;

(first entry) 26-MAR-2002 Colony stimulating factor 1 receptor (CSF1R) oligonucleotide #40.

Colony stimulating factor 1 receptor; CSF1R; polymorphic variant; cytostatic; gene therapy; malignant histiocytosis; isogene; myeloid malignancy; inflammatory disorder; transgenic animal; haplotype; genotype; human; allele specific oligonucleotide; ASO; probe; ss.

Homo sapiens.

WO200179225-A2.

25-OCT-2001

12-APR-2001; 2001WO-US012044.

12-APR-2000; 2000US-0196411P.

(GENA-) GENAISSANCE PHARM INC

Chew A, Choi JY, Koshy B;

WPI; 2002-075058/10.

Novel polymorphic variants of colony stimulating factor 1 receptor useful in studying expression and function of the protein, useful for screening candidate drugs to treat diseases e.g. inflammatory disorders.

Claim 15; Page 15; 164pp; English

The invention describes a novel isolated polymucleotide (I) comprising a sequence which is a polymorphic variant (PV) of a reference sequence for colony stimulating factor I receptor (CSFIR) gane, found on The polypeptide are useful for improving the discovery and development of drugs for treating diseases associated with CSFIR activity, e.g., and inflammatory discorders and the haplotypes can be used to validate CSFIR as a candidate target for treating a specific condition or disease predicted to be associated with CSFIR and and the haplotypes can be used to validate CSFIR gene of an individual can also the used in developing diagnostic tests and therapeutic treatments. (I) is useful in studying the expression and function of CSFIR, and in cypressing CSFIR protein for use in screening for candidate drugs to expressing CSFIR protein for use in screening for candidate arings to the variation on the biological activity of CSFIR. Antibodies are useful in a variety of diagnostic and prognostic formats and therapeutic useful in a variety of diagnostic and prognostic formats and therapeutic contents and compounds. A transgenic animal is useful in studying expression of the cSFIR protein, and for testing the efficacy of the transpeutic agents and compounds. Allele specific oligonucleotides (ASC) are useful as probes and primers, and for assaying a polymorphism in the transpect region. Without requiring any a priori knowledge of the phenotypic effect of any particular CSPIR or assaying a polymorphisms, ceffect of any particular CSPIR or described that are more likely to show efficacy in clinical trials. This sequence is an allele specific.

Sequence 15 BP; 2 A; 7 C; 1 G; 4 T; 0 U; 1 Other;

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Length 15;
Score 9.8; DB 1; Le
Pred. No. 1.2e+03;
0; Mismatches 2;
13.4%;
84.6%;
                                        933 CCTCCTCTTCATT 945
                                                         CCTCCTCSTCAGT 13
                      Conservative
           Best Local Similarity
Matches 11; Conserv
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RESULT 1337

BP.

ABS51924 standard; DNA; 15

ABS51924;

(first entry) 05-NOV-2002 Human FMO2 gene polymorphism detection ASO primer #45.

Human; flavin containing monooxygenase-2; FMO2; isogene; drugs targeting; drug toxicity; bone disorder; gene therapy; polymorphism; chromosome lq; allele-specific oligonucleotide; ASO; primer; ss.

Homo sapiens.

WO200253579-A2

11-JUL-2002.

18-DEC-2001; 2001WO-US049059.

29-DEC-2000; 2000US-0259062P

(GENA-) GENAISSANCE PHARM INC

Ä Parks Messer C, Lee HH, Sentivegna SC, Duda A, Kazemi A,

WPI; 2002-590627/63.

Novel genetic variants of Flavin Containing Monooxygenase 2 isogenes, useful for improving efficiency and reliability in drug development for treating developmental bone disorders.

Claim 15; Page 16; 140pp; English.

The present invention relates to a new polynucleotide which comprises flavin containing monooxygenase-2 (FMO2) isogenes. The invention is consciented for drugs that are useful for treating drug toxicity.

The methods of the invention are useful for improving the efficiency and reliability of several steps in the discovery and development of drugs of for treating diseases associated with FMO2 activity. The methods are also used by the pharmaceutical research scientist to validate FMO2 as a candidate target for treating a specific condition or disease predicted to be associated with FMO2 activity, e.g. drug toxicity, and in the compounds targeting FMO2. The methods are also useful for screening compounds targeting FMO2. The nucleic acid of the invention is useful in studying the expression and function of FMO2, and in expression is useful in studying the effect of the variation on the biological activity of FMO2 as well as on the binding affinity of candidate drugs targeting FMO2 is a well as on the binding affinity of candidate drugs targeting FMO2 is swell as on the binding affinity of candidate drugs targeting FMO2 is the treatment of drugs toxicity. The invention is useful for studying the expression of FMO2 and compounds for in vivo screening and testing of drugs targeted against FMO2 protein, and for testing the efficacy of therapeutic agents and compounds for treating the efficacy of therapeutic agents and compounds for treating the efficacy of therapeutic agents of incompounds for treating the efficacy of therapeutic agents of invention to detect polymorphisms in the human FMO2 gene located on

G; 7 T; 0 U; 1 Other; Sequence 15 BP; 2 A; 5 C; 0

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S.4 .

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Query Match
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Sequence 15 BP; 11 A; 0 C; 3 G; 0 T; 0 U; 1 Other;

13.4%; Score 9.8; DB 1; Length 15;

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Gaps

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84.68;

11; Conservative

Matches

Best Local Similarity

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The present invention relates to a new polymucleotide which comprises flavin containing monocxygenase-2 (FMC2) isogenes. The invention is useful in screening for drugs that are useful for invention is the methods of the invention are useful for improving the efficiency and reliability of several steps in the discovery and development of drugs for treating diseases associated with FMC2 activity. The methods are also used by the pharmaceutical research scientist to validate FMC2 as a candidate target for treating a specific condition or disease predicted to be associated with FMC2 activity, e.g. drug toxicity, and in the design of clinical trials for treating a specific condition of disease associated with FMC2 activity. The methods are also useful for screening compounds targeting FWC2. The nucleic acid of the invention is useful in studying the expression and function of FMC2, and in expression for screening for candidate drugs to treat diseases to related to FMC2 activity. It is also useful in studying the effect of the variation on the biological activity of FMC2 as well as on the binding affinity of candidate drugs targeting FMC2 is sugainst FMC2 for in vivo screening and testing of drugs targeted against FMC2 protein, and for testing the expression of FMC2 is ogenes in vivo, for in vivo screening and testing of drugs targeted against FMC2 protein, and for testing the efficacy of therapeutic agents and compounds for treating drug toxicity in a biological system. The present nucleic acid sequence represents an allele-specific condition continued to the human FMC2 gene located on chromosome Iquect to detect polymorphisms in the human FMC2 gene located on chromosome Iquect.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human, flavin containing monooxygenase-2, FMO2, isogene, drugs targeting, drug toxicity, bone disorder, gene therapy, polymorphism, chromosome 1q, allele-specific oligonucleotide, ASO, probe, ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Novel genetic variants of Flavin Containing Monooxygenase 2 isogenes, useful for improving efficiency and reliability in drug development for treating developmental bone disorders.
                                                                                       Gaps
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0
       Score 9.8; DB 1; Length 15;
Pred. No. 1.2e+03;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Messer C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human FMO2 gene polymorphism detection ASO probe #21.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Kazemi A,
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84.6%;
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Query Match
Best Local Similarity 84.6
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ABS51876
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AAS51876
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ACC ABS51876
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Human; F
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Human; F
XX
Human; F
XX
Homo sag
XX
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The present invention relates to a set of oligonucleotide probes and methods for detecting several different target polynucleotides. The set comparises a collection of different promiscuous probes each of which is capable of hybridising to a target sequence shared between at least two target polynucleotides where one target polynucleotide comprises at least toe target sequence that is shared with one or more other polynucleotides. A predefined combination of promiscuous probes is apable of hybridising to target sequences of at least one target polynucleotide, wherein said predefined combination of probes provide specificity of detection of that target polynucleotide. The probes of the invention are useful for detecting a number of different target polynucleotides using a programmable digital computer or for detecting an uncharacterised number of a polynucleotide family. The present sequence is an oligonucleotide probe used to detect potyvinus potato virus y (PVY) polymerase B motif target DNA in the method of the
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                                                                                                                                                                                                                                                                                                                           Promiscuous probe, target nucleic acid, detection, polymerase, B motif, potato virus Y_i PVY, ss.
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                Gaps
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Pred. No. 1.2e+03;
0; Mismatches 2; Indels
                Indels
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1.2e+03;
ches 3;
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Pred. No. 1.2e
L; Mismatches
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                                                                                                                                                                        BP.
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17-AUG-2000; 2000AU-00009483.
18-AUG-2000; 2000US-0226212P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         27-JUL-2001; 2001WO-AU000931.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (AUSU ) UNIV AUSTRALIAN NAT
                                                                       15 TTTTTCMTTCCTTT 1
                                                                                                                                                                    AAD30483 standard; DNA; 15
                                                                                                                                                                                                                                                            (first entry)
Best Local Similarity 73.3
Matches 11; Conservative
                                                   926 TITIAICCICCICT
                                                                                                                                                                                                                                            (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gibbs MJ, Gibbs AJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2002-206194/26.
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                                                                                                                                                                                                                                                                                                                                                                                  Potato virus Y.
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                                                                                                                                                                                                                                        07-AUG-2003
21-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        07-FEB-2002.
                                                                                                                                                                                                                                                                                                                                                                                                                                     variation
                                                                                                                                                                                                         AAD30483;
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                                                                                                                                         RESULT 1339
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13 TGCCCTTTTTCCC 1

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RESULT 1341
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Set of oligonucleotide probes for detecting different target polynucleotides, comprises a collection of different promiscuous probes each of which hybridizes to a target sequence shared between two target
                                                                                                                                                                                                                                                                                                    Promiscuous probe, target nucleic acid, detection, polymerase, B motif, potato virus Y_1 PVY; ss.
                                                                                                                                                                                                                                                                  Probe #3 used to detect potyvirus PVY polymerase B motif target DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         virus Y (PVY) polymerase B motif target DNA in the metho invention. (Updated on 07-AUG-2003 to correct OS field.)
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                                                                                                                                 AAD30478 standard; DNA; 15 BP
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17-AUG-2000; 2000AU-00009483.
18-AUG-2000; 2000US-0226212P.
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                                                                                                                                                                                                         (revised)
(first entry)
922 TGCCTTTTATCCC 934
                                  13 recerriratree 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 2002-206194/26.
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                                                                                                                                                                                                                                                                                                                                                              Potato virus Y.
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21-MAY-2002
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                                                                                                                                                                                                                                                                                                                                                                                                     Key
variation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gibbs MJ,
                                                                                                                                                                     AAD30478;
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                                                                                            RESULT 134
AAD30478/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New genetic variants of isolated N-acetylgalactosaminidase (NAGA), Alpha gene, useful for therapeutic purposes, for studying the expression and function of the polynucleotide, and for expressing NAGA protein.
                                                                                                                                                                                                                                            Human; PCR; primer; ss; gene therapy; N-acetylgalactosaminidase alpha; chromosome 22q13.2-q13.31; lysosomal glycohydrolase; screening; SNP; NAGA-related disease; single nucleotide polymorphism; haplotyping; NAGA;
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                                                                                                                                                                                 Human N-acetylgalactosaminidase (NAGA) alpha gene ASO primer 21.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Parks KE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 16; Page 13; 91pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  specific oligonucleotide primer
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   BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Koshy B,
ABT05329 standard; DNA; 15
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                                                                                                                         24-OCT-2002 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200194637-A1.
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                                                                                                                                                                                                                                                                                                                                                  genotyping.
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                                                                 ABT05329;
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Matches
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11; Conservative

Local Similarity

Best Loca Matches

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Gaps

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New haplotypes of human apolipoprotein C-IV gene, useful to diagnose and treat diseases associated with its activity such as hypertriglyceridemia.
            Apolipoprotein C-IV; APOC4; human; antilipaemic; haplotyping; hypertriglyceridaemia; allele-specific oligonucleotide; ASO; ss.
Apolipoprotein C-IV allele-specific oligonucleotide #31.
                                                                                                                                                                                   Claim 16; Page 13; 64pp; English.
                                                                                                          (GENA-) GENAISSANCE PHARM INC
                                                                                                                                                                                                                                                                                                                                                                                                          ABL51962 standard; DNA; 15 BP
                                                                              10-APR-2001; 2001WO-US011715.
                                                                                            11-APR-2000; 2000US-0195825P
                                                                                                                                Choi JY, Kliem SE,
                                                                                                                                               WPI; 2002-041284/05
                                                                                                                  (геен/) гее н н.
                                                  WO200177127-A2
                                    Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
                                                               18-OCT-2001
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The invention relates to haplotyping the apolipoprotein C-IV (APOC4) gene of an individual, comprising determining if the individual has one of the APOC4 haplotypes or haplotype pairs fully defined in the secification. Haplotyping the APOC4 gene of an individual, comprises determining the identity of the nucleotide at two or more polymorphic sites in one copy of the gene. The method also comprises identifying an association between a trait and a haplotype or haplotype pair of the APOC4 gene, comprising the trait with that of a reference population. A higher frequency in the trait population indicates the trait is associated with the haplotype. The polymucleotides and screened compounds are useful for developing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             treatment for diseases associated with APOC4 activity such as hypertriglyceridaemia. AAS95580-AAS95634 represent human apolipoprotein C-IV allele-specific oligonucleotides of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 15 BP; 1 A; 2 C; 1 G; 10 T; 0 U; 1 Other;
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1 TCTTTTTGTTTTAYC 15

11-JUL-2002 (first entry)

Human SLC18A2 allele specific oligonucleotide probe SEQ ID NO:10.

Human; solute carrier family 18 member 2; SLC18A2; vesicular monoamine; vesicular monoamine transporter; VWAT2; polymorphic site; SNP; single nucleotide polymorphism; antinifammatory; neuroleptic; haplotyping; genotyping; respiratory inflammatory disease; neuropsychiatric disorder; monoaminergic brain system; probe; ss.

Location/Qualifiers 8 Key misc_feature

components of the biological activity and instruction of the variation of controlled and sequence (SI) comprising soluble carrier family 18 (vesicular monoamine), member 2 (SLC18A2) isosgene selected from 49 isosgenes with regions of a sequence (SS) of 40023 bp (see ABL51954), and defined by a corresponding sequence (SS) of 40023 bp (see ABL51954), and defined by a corresponding set of polymorphisms whose locations and identities are given in the specification; or a sequence (S2) complementary to (S1). (I) has antinflammatory and neuroleptic activities, and can be used in gene theretory and neuroleptic activities, and can be used in gene the arguesian and function of SLC18A2, and in expressing the SLC18A2 gene in an individual. SLC18A2 is also known as the vesicular monoamine transporter (WMAT2). (I) is useful in studying the vestication of SLC18A2, and in expressing the SLC18A2 protein for candidate drugs to treat diseases controlly of SLC18A2, and in expressing the variation on the biological activity of SLC18A2 as well as on the binding affinity of controlled activity of SLC18A2 as well as on the binding affinity inflammatory diseases such as neuropsychiatric disorders involving monoaminergic brain systems. The present sequence represents an allele ö monoaminergic brain systems. The present sequence represents an allele specific oligonucleotide (ASO) probe for human SLC18A2, which is given in /*tag= a /note= "polymorphic site indicated by an ambiguity base" The present invention describes an isolated polynucleotide (I) having a Novel genetic variants of soluble carrier family 18 (vesicular monoamine), member 2 gene useful for screening drugs to treat diseases e.g. neuropsychiatric disorders involving monoaminergic brain systems. Gaps ö 13.4%; Score 9.8; DB 1; Length 15; 73.3%; Pred. No. 1.2e+03; ive 1; Mismatches 3; Indels Sequence 15 BP; 0 A; 5 C; 4 G; 5 T; 0 U; 1 Other; Sausker EA; Claim 17; Page 14; 183pp; English. Kliem SE, (GENA-) GENAISSANCE PHARM INC. 17-SEP-2001; 2001WO-US042217. 15-SEP-2000; 2000US-0232895P. 11; Conservative the present invention WPI; 2002-393942/42. Best Local Similarity WO200222652-A2 21-MAR-2002. Query Match Matches

Koshy B;

933 CCTCCTCTTCATTGG 947 15 CCTGCTCYTCTGTGG

RESULT 1344

ABK96294 standard; DNA; 15 BP. 24-SEP-2002 (first entry) ABK96294; ABK96294,

EDG1 gene allele-specific oligonucleotide #9.

BDG1; human; haplotyping; vascular developmental disorder; PCR; primer; endothelial differentiation sphingolipid G protein-coupled receptor 1;

Homo sapiens. BXXXXXXXXXXXXXXX

WO200244200-A2 06-JUN-2002

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Duda A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 1346
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   à
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to an isolated polynucleotide (I) encoding endothelial differentiation, sphingolipid G protein-coupled receptor I (BDG1) (II). Also described are methods for haplotyping or genotyping EDG1 gene of an individual by identifying single mucleotide polymorphisms (SNPs) of the gene. (II) is useful in screening for drugs targeting (II) that are useful for treating vascular developmental disorders. The methods are useful for improving the efficiency and reliability of several steps in the discovery and development of drugs for treating several steps in the discovery and development of drugs for treating used in pharmaceutical research to validate EDG1 as a candidate target for treating a specific condition or disease predicted to be associated with EDG1 activity. The methods as a candidate target for treating a specific condition of disease associated with EDG1 activity. The methods are also useful for screening compounds targeting EDG1. ARK96286-ARK96312 represent EDG1 gene allelespecific oligonucleotides, primer extension oligonucleotides and related
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
                                                                                                                                                                                                                                                                                                          Novel genetic variants of Endothelial Differentiation, Sphingolipid G
Protein-Coupled Receptor 1 isogenes, useful for improving efficiency and
reliability in drug development for treating vascular developmental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; ss; allele specific oligonucleotide; primer;
single nucleotide polymorphism; SNP; lipase endothelial isogene; LIPG;
drug screening; atherosclerosis; cardiovascular disorder;
LIPG haplotyping; LIPG genotyping.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 15 BP; 6 A; 0 C; 8 G; 0 T; 0 U; 1 Other;
                                                                                                                                                                                                     Shah N;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 14; Page 13; 68pp; English.
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                                                                                                                                        (GENA-) GENAISSANCE PHARM INC
                              03-DEC-2001; 2001WO-US046946.
                                                                                     01-DEC-2000; 2000US-0250606P.
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                                                                                                                                                                                                     Kazemi A,
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Best Local Similarity
Matches 11; Conserv
                                                                                                                                                                                                     Bieglecki KM,
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AC ABL9
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CW HUME
XW SING
XW SING
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XW LIPP
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The present sequence is an allele-specific oligonucleotide probe that was designed to detect a specific polymorphism in the human Duffy blood group (FY) gene (see ABL57150). The probe, and a probe of complementary sequence, belong to a set of probes (see ABL57151-66) that can be used in a kit for haplotyping or genofyping the FY gene of an individual. The probes provide good discrimination between the different FY gene
                                                                                                                                                                                                                 The invention comprises the DNA and amino acid sequence of the human lipase, endothelial (LIPG) isogene. Specifically, the invention relates to the discovery of 20 novel polymorphic sites within the LIPG gene. The LIPG coding sequence and protein are useful for screening drugs that can be used to treat atherosclerosis and other cardiovascular disorders. The LIPG coding sequence can also be used to haplotype and genotype the LIPG gene of an individual. The DNA sequences ABL91822 - ABL91861 represent LIPG gene allele specific oligonucleotide primers
                                                                                   Novel genetic variants of Lipase, Endothelial isogenes, useful for improving efficiency and reliability in drug development for treating diseases associated with LIPG activity, e.g. atherosclerosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Duffy; blood group; FY; human; receptor; haplotyping; genotyping; transgenic animal; malaria; inflammation; antimalarial; protozoacide; antiinflammatory; single nucleotide polymorphism; SNP; probe; ss.
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73.3%; Pred. No. 1.2e+03;
ive 1; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 15 BP; 0 A; 4 C; 1 G; 9 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Probe for FY gene polymorphism detection.
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  Messer
                                                                                                                                                                             Claim 16; Page 14; 134pp; English.
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Kliem SE,
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Best Local Similarity 73.3%
Matches 11, Conservative
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                                             WPI; 2002-292055/33
  Kazemi A,
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Gaps

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The invention relates to a typing kit for judging human leukocyte antigen (HLA) genotype of a sample by hybridising a substrate on which 10-24 base oligonucleotides (ABLSOS12-ABLSISES) originating in the sequences of genes e.g. belonging to HLA class I antigens on human genome and containing gene polymorphisms as alloantigens have been immobilised as primers for amplification of cleaved nucleic acids relating to gene polymorphisms. The method is useful for judging HLA genotypes of individuals by determining immunogenetic differences before transplanting between them, providing genetic information to decide compatibility of organ and tissue for transplantation e.g. of bone marrow, kidney, liver, pancreas, Langerhans islet in pancreas and cornea, susceptibility diagnosis of genetic diseases and identifying individuals
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human leukocyte antigen (HLA) typing, useful for judging HLA genotypes of individuals e.g. by determining immunogenetic differences when transplanting between them.
polymorphisms by each having a central nucleotide that aligns with the polymorphic site in the target region. The present invention provides novel genetic variants of the FY gene, and discloses various genotypes, haplotypes and haplotype pairs that exist in the general United States population. Compositions and methods for haplotyping and/or genotyping the FY gene in an individual are also disclosed. The polymorphism and haplotype data are useful for validating FY as a candidate target for treating a condition or disease associated with FY activity, such as malaria and inflammatory disorders
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; human leukocyte antigen; HLA; genotype; polymorphism; immunogenetic; transplantation; genetic disease; ss.
                                                                                                                                                                                                                                         13.4%; Score 9.8; DB 1; Length 15; 73.3%; Pred. No. 1.2e+03; ative 1; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human HLA genotyping oligonucleotide SEQ ID NO 24.
                                                                                                                                                                                                     Sequence 15 BP; 0 A; 5 C; 0 G; 9 T; 0 U; 1 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 10; Page 97; 345pp; Japanese.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-JUN-2001; 2001WO-JP004662.
                                                                                                                                                                                                                                                                                                                       921 TIGCCTTTTATCCCT 935
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Best Local Similarity
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13.4%; Score 9.8; DB 1; Length 15;

Query Match

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; PCDH2; protocadherin 2; haplotyping; polymorphic variant; SNP;
               Gaps
                                                                                                                                                                                                                                              Human, colon cancer; colorectal cancer; pancreatic cancer; SAGE tag; serial analysis of gene expression; diagnostic; prognostic; probe; cancer marker; ss.
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               Indels
  Pred. No. 1.2e+03;
; Mismatches 2;
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84.68;
                                                                                                                                                                                                                        Human colon cancer SAGE tag
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                                           931 TCCCTCCTCTTCA 943
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  Best Local Similarity 84.6
Matches 11; Conservative
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The invention relates to haplotyping the protocadherin 2 (PCDH2) gene, comprising determining which of the haplotypes given in the specification defines one or both opies of the individual's gcDH2 gene. The polymorphiems are within a 30244 base pair sequence (ABAD5413). Fully defined in the specification. The polymorphic variants are useful in studying the expression and function of PCDH2, in expressing PCDH2 protein for use in screening for candidate drugs to treat diseases such as cancer, related to PCDH2 activity, in studying the effect of the variation on the biological activity of PCDH2 and the binding affinity of candidate drugs targeting PCDH2. The haplotyping methods are useful in validating PCDH2 as a candidate target for treating a specific condition or disease predicted to be associated with PCDH2 activity or in the design of clinical trials of candidate drugs for treating a specific condition or disease associated with PCDH2 activity. The present sequence is that of a PCDH2 allele-specific oligonucleotide (ASO) PCR primer of
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                                                                                                                                                                                                                                                                                                                                    New protocadherin 2 (PCDH2) polymorphic variants and encoding genes, useful in expressing PCDH2 protein for screening candidate drugs to treat diseases related to PCDH2 activity.
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single nucleotide polymorphism; cytostatic; cancer; chromosome 5q31; allele-specific oligonucleotide; ASO; PCR primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                         Claim 16; Page 13; 127pp; English.
                                                                                                                                                                                                                                                               Tanguay DA;
                                                                                                                                                                                                                               (GENA-) GENAISSANCE PHARM INC.
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                                                                                                                                                                                                                                                                 Koshy B,
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                                                                                     WO200194361-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                the invention
                                                      Homo sapiens
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                                                                                                                         13-DEC-2001.
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Best Local S
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The invention relates to haplotyping the protocadherin 2 (PCDH2) gene, comprising determining which of the haplotypes given in the specification defines one or both ocopies of the individual's pCDH2 gene. The polymorphiems are within a 30344 base pair sequence (ABA05413), fully defined in the specification. The polymorphic variants are useful in constitution to the expression and function of PCDH2, in expressing PCDH2 controling for candidate drugs to treat diseases such as cancer, related to PCDH2 activity, in studying the effect of the variation on the biological activity of PCDH2 and the binding affinity of candidate drugs trageting PCDH2. The haplotyping methods are useful in validating PCDH2 as a candidate target for treating a specific condition or disease associated with PCDH2 activity or in the condition or disease associated with PCDH2 activity. The present sequence condition or disease associated with PCDH2 activity. The present sequence is that of a PCDH2 allele-specific oligonucleotide (ASO) PCR primer of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                    New protocadherin 2 (PCDH2) polymorphic variants and encoding genes, useful in expressing PCDH2 protein for screening candidate drugs to diseases related to PCDH2 activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human, neuropeptide Y, NPY, isogene, SNP, atherosclerosis, obesity, psychological disorder, single nucleotide polymorphism; alcoholism; antiarteriosclerotic; anorectic; PCR, primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 13.4%; Score 9.8; DB 1; Length 15; 73.3%; Pred. No. 1.2e+03; tive 1; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human neuropeptide Y allele specific primer SEQ ID NO: 40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Stephens JC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 15 BP; 8 A; 2 C; 3 G; 1 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Nandabalan K,
                                                                                                                                                                                                                                         Claim 16; Page 14; 127pp; English.
                                                                                                   Tanguay DA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Lanz EM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              BP.
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                                                                    (GENA-) GENAISSANCE PHARM INC.
06-JUN-2001; 2001WO-US018321.
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                                  06-JUN-2000; 2000US-0209564P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAL48116 standard; DNA; 15
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                                                                                                   Koshy B,
                                                                                                                                    WPI; 2002-097928/13
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                                                                                                     Kliem SE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAL48116;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 1351
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Matches
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The present invention provides the human neuropeptide Y (NPY) gene and single nuclectide polymorphisms (SNPs) identified therein. The sequence can be used in the treatment of disorders associated with NPY, including atherosclerosis, obesity, psychological disorders and alcoholism. The present sequence is an allele specific primer used to isolate the human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present invention provides the human neuropeptide Y (NPY) gene and single nucleotide polymorphisms (SNPs) identified therein. The sequence can be used in the treatment of disorders associated with NPY, including atherosclerosis, obesity, psychological disorders and alcoholism. The present sequence is an allele specific primer used to isolate the human
         New genetic variants of the human Neuropeptide Y (NPY) gene useful for treating disorders affected by abnormal expression or function of NPY isogene e.g., atherosclerosis or obesity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New genetic variants of the human Neuropeptide Y (NPY) gene useful fortreating disorders affected by abnormal expression or function of NPY
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                                                                                                                                                                                                                                                                                            Query Match 13.4%; Score 9.8; DB 1; Length 15; Best Local Similarity 84.6%; Pred. No. 1.2e+03; Matches 11; Conservative · 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human neuropeptide Y allele specific primer SEQ ID NO: 42.
                                                                                                                                                                                                                                                        Sequence 15 BP; 7 A; 0 C; 7 G; 1 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 11; Page 17; 80pp; English.
                                                                                        Claim 11; Page 17; 80pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (GENA-) GENAISSANCE PHARM INC
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                                                                                                                                                                                                                                                                                                                                                                       925 CTTTTATCCCTCC 937
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAL48118 standard; DNA; 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              27-SEP-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                            CTTTTCTCCCTTC
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                                                                                                                                                                                                                       NPY coding sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 1352
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Stephens JC;

Nandabalan K,

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The invention relates to a novel pharmaceutical composition, which has a first active agent comprising an oligonucleotide antisense to the intitation codon, coding region, 5' or 3' end genomic flanking regions, 5' and 3' intron-exon junctions, or regions within 2-10 nucleotides of junctions of genes encoding a polypeptide associated with lung and/or nasal airway dysfunction and a second active agent comprising an antiinflammatory steroid and ubiquinone. A composition of the invention has antiinflammatory, antiallergic, antistachmatic, hypotensive, immunosuppressive, and cytostatic activity. The composition may have a preventing a respiratory, lung or malignant disease or condition, also for enhancing the prophylactic or therapeutic respiratory effect of an preventing sensitivity to adenosine, reducing levels of adenosine receptor, producing bronchodilation, increasing levels of ubiquinone or lung inflammation, lung allergies, or a respiratory disease or condition, lung inflammation, but was obtained in electronic format directly from WIPO are fear into into into into the printed of the first into into into into the printed of the first into into into into increasing format directly from WIPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pharmaceutical composition for treating ailments associated with impaired respiration, has oligo(s) antisense to specific gene(s) or its corresponding RNAs, and glucocorticoid or non-glucocorticoid steroid or
                                                                                                                                                                                                                                                                                          Human; antisense; lung dysfunction; nasal airway dysfunction; antiinflammatory steroid; ubiquinone; antiinflammatory; antiallergic; antiasthmatic; hypotensive; cytostatic; gene therapy; antisense gene therapy; respiratory; lung; adenosine sensitivity; adenosine receptor; bronchodilation; bronchoconstriction; lung allergy; lung inflammation; respiratory disease; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pabalan J, Aguilar D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 15 BP; 0 A; 5 C; 2 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                          Human IL3 receptor antisense fragment no.1094.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; SEQ ID NO 10472; 872pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            at ftp.wipo.int/pub/published_pct_sequences
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S;
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Tang L, Shahabuddin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   24-APR-2001; 2001US-0286137P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (EPIG-) EPIGENESIS PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             23-APR-2002; 2002WO-US013135
                                                                                                                                     ABZ95230 standard; DNA; 15
CITITATCCCTCC 937
                                                                                                                                                                                                                    17-0CT-2003 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
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Miller S,
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925
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Gaps

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Gaps

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Query Match 13.4%; Score 9.8; DB 1; Length 15; Best Local Similarity 84.6%; Pred. No. 1.2e+03; Matches 11; Conservative 0; Mismatches 2; Indels

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Gaps

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Query Match
13.4%; Score 9.8; DB 1; Length 15;
Best Local Similarity 84.6%; Pred. No. 1.2e+03;
Matches 11; Conservative 0; Mismatches 2; Indels

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Cell typing, intergene region, IGR1; IGR2; external transcribed space, ETS1; ETS2; unknown cell type, prokaryotic cell; bacterium; eukaryotic cell; bacterial strain identification, nosocomial infection; species identification; dedicree; prokaryotic cellsm; identification of pathogenic bacteria; l6S ribosomal RNA; 16S rRNA; PCR; primer; ss.
                                                                                                Universal PCR primer B-C for prokaryotic 168 rRNA gene.
                                                                                                                                                                                                                   01-NOV-2001; 2001US-00001048.
                                                                                                                                                                                                                                   95US-00461210.
                                                                                                                                                                                                                                                                               Whitehouse E,
                                             .120/c
ABX11120 standard; DNA; 15
938 TCTTCATTGGTTT 950
                                                                                  (first entry)
        TCTTCCTTCGTTT 13
                                                                                                                                                                                                                                                 (LEGG/) LEGGETT C G.
(WHIT/) WHITEHOUSE E.
(REEV/) REEVES R H.
                                                                                                                                                                                    US2002164610-A1.
                                                                                                                                                                      Bacteriaceae
                                                                                                                                                                                                                                   05-JUN-1995;
                                                                                  28-APR-2003
                                                                                                                                                                                                                                                                                Leggett CG,
                                                                                                                                                                                                    07-NOV-2002.
                                                                  ABX11120;
                                      RESULT 1354
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Typing cell by amplifying nucleic acid from cells of unknown type having nucleic acid forming portion of cell's genome, digesting and separating product to generate pattern, comparing pattern with known cell pattern. WPI; 2003-247254/24.

Disclosure; Page 3; 24pp; English.

Reeves

New genetic variants having polymorphisms in the G protein, GNB3 gene, useful for treating disorders with abnormal expression or function of the GNB3 gene, such as asthma, obesity, hypertension and left ventricular

Claim 16; Page 14; 88pp; English.

hypertrophy.

The present invention relates to a method for typing a cell. The method comprises (a) providing a cell of unknown type having a nucleic acid comming a portion of the cell's genome such as intergene region (ISR1, ISR2, or external transcribed space (ETS) or ETS2, (b) isolating the connected space (ETS) or ETS2, (b) isolating the nucleic acid NA from the unknown cell type, amplifying the nucleic acid NA from the unknown cell type, amplifying the nucleic acid NA from the unknown cell type, amplifying the nucleic acid the restriction pattern, and comparing the restriction pattern, and comparing the restriction pattern of a cell of known type. The method is useful for the restriction pattern of a cell of known type. The method is useful for identifying differences between the individuals of the subspecies, subspecies of the same species, and for identifying species, subspecies of the method is useful for identifying differences between the individuals of the subspecies of higher life forms. The method of the individuals of subspecies of higher life forms. The method of the currently practiced methodologies, allows the quantitative analysis of the subspecies of hubber as few hours, as opposed to the lengthy turnarcound time associated with the methods of Southern hybridisation and Dot Blots, and allows rapid strain identification of pathogenic backeria. ABXIIII77 and allows repaid strain identification of pathogenic backeryotic I6S RNA (rRNA) genes

Sequence 15 BP; 6 A; 2 C; 5 G; 2 T; 0 U; 0 Other;

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                                                                                                                                                           Human, guanine nucleotide binding protein beta polypeptide 3, G protein, GNB3, polymorphism; obesity, left ventricular hypertrophy, hypertension; drug discovery; cardiovascular; development process; asthma; anorectic;
                 Gaps
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Length 15;
Score 9.8; DB 1; Length 15
Pred. No. 1.2e+03;
0; Mismatches 2; Indels
                                                                                                                                            Human GNB3 gene polymorphisms detecting ASO probe #16.
                                                                                                                                                                                                                                                                                                          Koshy
                                                                                                                                                                                                                                                                                                         Kliem SE,
                                                                                            BP.
                                                                                                                                                                                                                                                                                         (GENA-) GENAISSANCE PHARM INC.
                                                                                                                                                                                                                                                         21-MAR-2001; 2001WO-US008961.
                                                                                                                                                                                                                                                                         21-MAR-2001; 2001WO-US008961.
 13.4%;
84.6%;
                                                                                            AAD47748 standard; DNA; 15
                                  904 GICATITICITIG 916
                                                                                                                            (first entry)
Query Match 13.4
Best Local Similarity 84.6
Matches 11; Conservative
                                                                                                                                                                                                                                                                                                         Choi JY,
                                           13 GTCAATTCCTTTG 1
                                                                                                                                                                              drug discovery; cardiov;
gene therapy; probe; ss
                                                                                                                                                                                                                                                                                                                         WPI; 2003-018947/01.
                                                                                                                                                                                                                      WO200277284-A1.
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                                                                                                                            24-FEB-2003
                                                                                                                                                                                                                                        03-OCT-2002
                                                                                                           AAD47748;
                                                                           RESULT 1355
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The invention relates to an isolated polypeptide which comprises a first nucleotide sequence which is a polymorphic variant of a reference sequence for the guamine nucleotide binding protein (G protein), beta polymorphic variants of the GNB3 gene or fragment. Polymorphic variants of the GNB3 gene are useful in studying the expression and biological function of GNB3 and in identifying drugs targetting GNB3 protein for treating disorders associated with abnormal expression or function of GNB3, e.g. hypertension, obesity, asthma and left ventricular hypertrophy.

CC Polymorleotides comprising a polymorphic gene variant or fragment may be used for therapeutic purposes, where a patient could benefit from expression or increased expression of a particular GNB3 gene isoform or an expression vector encoding the isoform may be administered to the patient. Haplotype information is useful in improving the efficiency and cutput of several steps in drug discovery and development process, including target validation, identifying lead compounds and early phase clinical trials. The invention is used in gene therapy. The present constitution is a used in gene therapy. The present constitution is used in gene therapy. The present constitution is a used in gene therapy. The present constitution is a used in gene therapy. ô Gaps ö Length 15; 13.4%; Score 9.8; DB 1; Length 15 73.3%; Pred. No. 1.2e+03; tive 1; Mismatches 3; Indels Sequence 15 BP; 0 A; 5 C; 0 G; 9 T; 0 U; 1 Other; Query Match Best Local Similarity 73.35 Matches 11; Conservative oligonucleotide) probe

924 CCTTTTATCCCTCCT 938

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1 CTTTTTTWCTCTCCT = = = = =

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The invention describes an enzymatic nucleic acid molecule (1) which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B (NFRS), where (1) is an inozyme, zinzyme, G-cleaver or amberzyme configuration. The enzymatic nucleic acid molecule is adapted to treat cancer and is useful for down-regulating REL-A activity in a cell, for treating a parient having a condition associated with the level of REL-A. (1) is useful for cleaving RNA comprising a sequence of REL-A gene, in the presence of a divalent cation, especially MG^2+. The enzymatic and the presence of a divalent cation, especially MG^2+. The enzymatic and antisense nucleic acid molecules are useful for treating breast, lung, prostate, colorectal brain, oesophageal, stomach, bladder, pancreatic, cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or multidrug resistant cancer. The method involves use of other drug cervical, head and neck, ovarian cancer, melanoma, lymphoma, glioma or multidrug resistant cancer. The method involves use of other drug chemocherapy including paclitaxal, docetaxel, cisplatin, methotrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, edatrexate, cyclophosphamide, doxorubin, fluorouracil carboplatin, edatrexate, gencilabine or radiation therapy. The enzymatic and antisense nucleic caid molecules are also useful for treating inflammatory disease such as rheumatoid arthritis, restenosis, asthma, Crohn's disease, diabetes,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Novel enzymatic nucleic acid molecules which down regulates expression of a sequence encoding a subunit of nuclear factor kappa B useful for treating cancer, inflammatory disorders and autoimmune diseases.
                                                                                                                                                                                    Enzymatic nucleic acid; nuclear factor kappa B; NFKB; inozyme; zinzyme; G-cleaver; amberzyme; cancer; REL-A activity, breast cancer; human; lung cancer; prostate cancer; colorectal cancer; brain cancer; colorectal cancer; prostate cancer; colorectal cancer; pancatic cancer; cervical cancer; pand and neck cancer; voraian cancer; melanoma; lymphoma; glioma; multidrug resistant cancer; REL-A-specific inhibitor; chemotherapy; paclitaxel; docetaxel; cisplatin; methotrexate; cyclophosphanide; doxorubin; fluctouracil carboplatin; edatrexate; gencitabine; radiation therapy; inflammatory disease; asthma; diabetes; rheumatoid arthritis; restenosis; Crohn's disease; obesity; ischaemia; gene therapy; autoimmune disease; lupus; multiple sclerosis; sepsis; transplant/graft rejection; reperfusion injury; glomerulonephritis; allergic airway inflammatory bowel disease; infection; ss.
                                                                                                                                            Necrosis factor kappa B sub-unit modulating enzyme target #132
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Draper KG;
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94US-00245466.
94US-00291932.
96US-00777916.
ACA09939 standard; RNA; 15 BP
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                                                                                              03-JUN-2003 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US2002177568-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
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18-MAY-1994;
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                                                ACA0939;
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obesity, autoimmune disease, lupus, multiple sclerosis, transplant/graft rejection, gene therapy applications, ischaemia/repetitusion injury (central nervous system (CNS) and myocardial), glomerulonephritis, sepsis, allergic airway inflammation, inflammatory bowel disease or
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                                                                                                                                                                                                                                                                                                                                                                                                             Nucleic acid molecule, Hepatitis C virus, HCV; Hepatitis B virus; HBV, RNA stability; RNA expression; RNA synthesis; antisense; enzymatic nucleic acid; hammerhead ribozyme; DNAzyme; inozyme; zinzyme;
                                                    infection. This sequence represents the substrate of a novel enzymatic nucleic acid molecule
                                                                                                                                                                                                                                                                                                                                                                                                                                                       amberzyme, G-cleaver ribozyme; decoy molecule; aptamer; thosamer amberzyme; G-cleaver ribozyme; decoy molecule; aptamer; they reverse transcriptase; Enhancer I region; viral replication; degenerative; disease state; HBV infection; HCV infection; cirrhosis; liver failure; hepatocellular carcinoma; hepatotropic; cytostatic; virucide; antiinflammatory; substrate; ss.
                                                                                                                                                           Gaps
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                                                                                                                              13.4%; Score 9.8; DB 1; Length 15; 84.6%; Pred. No. 1.2e+03; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                    HBV enzymatic nucleic acid substrate sequence #88.
                                                                                                   Sequence 15 BP; 5 A; 3 C; 6 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mcswiggen J, Morrissey D,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Example 1; Page 214; 387pp; English.
                                                                                                                                                                                                                                                                                              ACD56199 standard; RNA; 15 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              26-MAR-2001; 2001US-00817879.
08-UN-2001; 2001US-00877478.
08-UN-2001; 2001US-0296876P.
24-OCT-2001; 2001US-0337055P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   26-MAR-2002; 2002WO-US009187.
                                                                                                                                               84.6%;
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Roberts E;
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MCSWIGGEN J.
MORRISSEY D.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2003-229207/22
                                                                                                                                               Best Local Similarity
Matches 11; Conser
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DRAPER K.
ROBERTS E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hepatitis B virus
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Draper K,
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                                                                                                                                                                                                                                                                                                                             ACD56199;
                                                                                                                                  Query Match
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(BLAT/)
(MACE/)
(MCSW/)
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(PAVC/)
(LEEP/)
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(ROBE/)
                                                                                                                                                                                                                                                                RESULT 1357
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The present invention relates to nucleic acid molecules which modulate

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Mon Oct 18 14:40:13 2004
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schultzi-899.rng

Novel compound useful for treating cirrhosis, liver failure, hepatcocallular carcinoma, or condition associated with hepatitis C virus

WPI; 2003-229207/22

infection.

the synthesis, expression and/or stability of Hepatitis C virus (HCV) or Hepatitis B virus (HBV) RNA. The nucleic acid molecules include antisense and antymatic nucleic acids such as hameerhead riboxymes. DNAzymes, inoxymes, amberzymes, and G-cleaver riboxymes. Also disclosed are nucleic acid decoy molecules and g-cleaver riboxymes. Also disclosed transcriptase and/or HBV reverse transcriptase primer sequences, as well as oligonucleotides that specifically bind the Enhancer I region of HBV DNA. The nucleic acids may be used to modulate the expression of HBV genes and HBV viral replication. Also disclosed is a method for screening compounds and/or potential therapies directed against HBV, and compounds that modulate the expression and/or replication of HCV. The compounds that modulate the expression and HCV infection, replication and gene actives a states related to HBV and HCV infection, replication and gene discase states related to HBV and HCV infection, replication and gene carcinoma. The present sequence represents a substrate for one of the HBV enzymatic nucleic acid sequence disclosed in the present invention Nucleic acid molecule, Hepatitis C virus, HCV, Hepatitis B virus, HBV, RNA stability, RNA expression, RNA synthesis, antisense, enzymatic nucleic acid, hammerhead ribozyme; DNazyme; inozyme; amberzyme; G-cleaver ribozyme; decoy molecule; aptamer; HBV reverse transcriptase; Enhancer I region, viral replication, degenerative, disease state, HBV infection; HCV infection; cirrhosis; liver failure; hepatocellular carcinoma; hepatotropic; cytostatic; virucide; antiinflammatory; substrate; ss. 0; Gaps Query Match
13.4%; Score 9.8; DB 1; Length 15;
Best Local Similarity 38.5%; Pred. No. 1.2e+03;
Matches 5; Conservative 6; Mismatches 2; Indels HBV enzymatic nucleic acid substrate sequence #150. Sequence 15 BP; 2 A; 5 C; 2 G; 0 T; 6 U; 0 Other; 26-MAR-2001, 2001US-00817879. 08-JUN-2001, 2001US-00877478. 08-JUN-2001, 2001US-0296876P. 24-OCT-2001, 2001US-0335059P. 05-DEC-2001, 2001US-0337055P. 26-MAR-2002; 2002WO-US009187 RIBOZYME PHARM INC. BLAIT L. ACD56425 standard; RNA; 15 929 TATCCCTCCTTT 941 (first entry) 3 UAUGCCUCAUCUU 15 MACEJAK D.
MCSWIGGEN J.
MORRISSEY D.
PAVCO P.
LEE P.
DRAPER K.
ROBERTS E. Hepatitis B virus WO200281494-A1. 24-SEP-2003 17-CCT-2002 ACD56425; (RIBO-)
(BLAT/)
(MACE/)
(MCSW/)
(MORR/)
(PAVC/)
(LEEP/)
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(ROBE/) ACD56425
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The present invention relates to nucleic acid molecules which modulate the synthesis, expression and/or stability of Hepatitis C virus (HCV) or Hepatitis B virus (HBN) RNA. The nucleic acid molecules include antisense and enzymatic nucleic acids such as hammerhead ribozymes, DNAzymes, inozymes, zinzymes, amberzymes, and G-cleaver ribozymes. Also disclosed transcriptase and/or HBV reverse transcriptase primer sequences, as well or soligonucleotides that specifically bind the Enhancer I region of HBV DNA. The nucleic acids may be used to modulate the expression of HBV CC genes and HBV viral replication. Also disclosed is a method for screening compounds and/or potential therapies directed against HBV, and compounds that modulate the expression and/or replication of HCV. The compounds of the invention are useful for the treatment of degenerative and disease states related to HBV and HCV infection, replication and gene expression such as cirrhosis, liver failure, and hepatocellular carcinoma. The present sequence represents a substrate for one of the HBV compounds.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human; cytochrome P450 subfamily IIE; CYP2E protein; haplotyping;
genotyping; gene therapy; cancer; allele-specific oligonucleotide; ASO;
polymorphism; probe; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New genetic variants comprising haplotypes of the cytochrome P450, subfamily IIE (CYP2E) gene, useful for screening drugs for treating cancer, validating CYP2E protein as a drug target, or reducing bias clinical trials of such drugs.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             13.4%; Score 9.8; DB 1; Length 15; 38.5%; Pred. No. 1.2e+03; tive 6; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sausker EA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human CYP2E gene polymorphism detecting ASO probe #4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 15 BP; 0 A; 6 C; 3 G; 0 T; 6 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Koshy B,
                                                                                                                  Example 1; Page 219; 387pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A, Gilson CR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAD51625 standard; DNA; 15 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   07-MAY-2002; 2002WO-US014540.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    917 GTCTTTGCCTTTT 929
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Best Local Similarity 38.5
Matches 5; Conservative
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AAD51625
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Д, Pavco

Morrissey D,

Mcswiggen J,

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Macejak D, Roberts I

Blatt L, ! Draper K,

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The invention relates to genetic variants of human cytochrome P450, subfamily IIE (CYPZE) gene. The invention also relates to compositions and methods for haplotyping and/or genotyping the CYPZE gene in an individual. The polynucleotide comprising polymorphisms in the CYPZE are useful in screening candidate drugs to treat diseases related to CYPZE activity, e.g. cancer. The methods and haplotypes are useful in improving the efficiency of drug discovery and development processes, or for designing clinical trials of candidate drugs for treating the specific condition or disease. The polymorphisms and haplotypes of CYPZE gene are useful for validating whether CYPZE is a suitable target for specific condition or discoase. The polymorphisms and haplotypes of CYPZE gene are useful for validating whether CYPZE is a suitable target for synthesis in cells, screening for drugs and reducing bias in clinical trials of the drugs. The invention is also useful in gene therapy. The present squares is an allele-specific oligomucleotide (ASO) probe used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
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Claim 37; Page 15; 94pp; English
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Matches
\overset{\alpha}{\times}\overset{\times}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overset{\circ}{\times}\overse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ABF75857 standard; DNA; 13 22-FEB-2002 (first entry) ABF75857; RESULT 1360

Oligonucleotide SEQ ID NO 175854 for detecting SNP TSC0043670.

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Homo sapiens

WO200177384-A2.

18-OCT-2001

06-APR-2001; 2001WO-IB000713.

07-APR-2000; 2000DE-01019173

(EPIG-) EPIGENOMICS

oligonucleotides, useful for diagnosis and cell typing, ied to detect single-nucleotide polymorphisms and cytosine Berlin K; Olek A, Piepenbrock C, WPI; 2001-657177/75. Set of oldesigned

13

Claim 1; SEQ ID NO 175854; 29pp + Sequence Listing; German.

methylation status.

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99889, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABH82073

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Gaps

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Ouery Match 13.2%; Score 9.6; DB 1; Length 13; Best Local Similarity 90.0%; Pred. No. 1.2e+03; Matches 9; Conservative 1; Mismatches 0; Indels

949 TTAATGTATC 958

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Sequence 13 BP; 4 A; 0 C; 3 G; 5 T; 0 U; 1 Other;

ftp.wipo.int/pub/published_pct_sequences

ö This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, contral nervous system, cardiovascular and metebolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF9989, ABH0010-ABH99989 and ABI00010-ABI82073 tepresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic. Gaps Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status. ; 0 Oligonucleotide SEQ 1D NO 175853 for detecting SNP TSC0043670. 13.2%; Score 9.6; DB 1; Length 13; 90.0%; Pred. No. 1.2e+03; ive 1; Mismatches 0; Indels Claim 1; SEQ ID NO 175853; 29pp + Sequence Listing; German Sequence 13 BP; 5 A; 3 C; 0 G; 4 T; 0 U; 1 Other; Berlin K; ABF75856 standard; DNA; 13 BP. 06-APR-2001; 2001WO-IB000713. 07-APR-2000; 2000DE-01019173 22-FEB-2002 (first entry) Local Similarity 90.0 les 9, Conservative 949 TTAATGTATC 958 Piepenbrock C, (EPIG-) EPIGENOMICS AG 10 TTAATGTATY 1 WPI; 2001-657177/75. WO200177384-A2 Homo sapiens. ABF75856; Query Match olek A, RESULT 1361 Best Loca Matches ABF75856 88866888 g ઠ schultz1-899.rng

ABF82142; RESULT 1364 à ď ö This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretracted genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for Aberoular and metabolic disorders. The ABC9989, ABF00010-ABF99899 and ABI00010-ABF80013 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at the printed specification, but fire wipo.int/pub/published_pct_sequences SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardicvascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic. 0; Gaps bet or oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status. Oligonucleotide SEQ ID NO 23516 for detecting SNP TSC0005018. 13.2%; Score 9.6; DB 1; Length 13; 90.0%; Pred. No. 1.2e+03; tive 1; Mismatches 0; Indels Claim 1; SEQ ID NO 23516; 29pp + Sequence Listing; German. Sequence 13 BP; 7 A; 2 C; 0 G; 3 T; 0 U; 1 Other; Berlin K; ABC23499 standard; DNA; 13 BP 07-APR-2000; 2000DE-01019173 06-APR-2001; 2001WO-IB000713 (first entry) Query Match
Best Local Similarity 90.v Olek A, Piepenbrock C, TTAATGTATC 958 (EPIG-) EPIGENOMICS AG |||||||||| TTAATGTATY 1 WPI; 2001-657177/75. TTAATGTATY WO200177384-A2 20-FEB-2002 Homo sapiens 18-OCT-2001. 949 10 ABC23499 RESULT 1363 à q d

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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Pred. No. 1.2e+03;
1; Mismatches 0; Indels
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07-APR-2000; 2000DE-01019173.

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Oligonucleotide SEQ ID NO 182140 for detecting SNP TSC0045025.

or peptide nucleic

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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Best Local Similarity
Matches 9; Conserv
                                                                          methylation status
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acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomucleotides are used for diagnosis and/or prognosis of cancer and range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH0010-ABF99989, ABH0010-ABF99989 and ABI0010-ABI32073 data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at
This invention describes novel oligonucleotide primers
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Best Local 9, Conservative
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                                                                                                                                                                                                  Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine
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Pred. No. 1.2e+03;
1; Mismatches 0; Indels
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Score 9.6; DB 1; Length 13; Pred. No. 1.2e+03; 1; Mismatches 0; Indels

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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nuclectide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic discorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABC0010-ABC99989, ABC0010-ABC99989, ABC0010-ABC99989 ABC0010-ABC99989 and ABI0010-ABC8030 represent the oligomers described in the invention. NOTE: The sequence was obtained in electronic format from WIPO at the printed specification, but the wipo.int/pub/published_pct_sequences
                                                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                            Oligonucleotide SEQ ID NO 23515 for detecting SNP TSC0005018.
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ABC23498 standard; DNA; 13 BP.
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Claim 1; SEQ ID NO 170273; 29pp + Sequence Listing; German.

Berlin

BP.

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New mammalian model for enhanced wound healing - useful for identifying
                                                                                                       Wound healing, non-MRL healer mouse, quantitative trait locus, QTL, healing response, microsatellite marker; treatment, central nerve; peripheral nerve, nerve injury; SAGE tag; murine; ss.
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                                                                               Murine C57BL/6 SAGE tag 1568982.
                                                                                                                                                                                                                                                                                                                                                                                                                          enhanced wound healing genes.
AAZ18749 standard; DNA; 11
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                                                                                                                                                                                                                                                                                 SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                         13.2%; Score 9.6; DB 1; Length 13; 90.0%; Pred. No. 1.2e+03; ive 1; Mismatches 0; Indels
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 Seguence 13 BP; 3 A; 0 C; 2 G; 7 T; 0 U; 1 Other;
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Matches 9; Conservative
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98US-0074737P. 98US-0097937P. 98US-0102051P.

99WO-US002962

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This invention describes a novel non-MRL healer mouse (M) having at least one quantitative trait locus selected from those given in the greification, exhibiting an enhanced healing response to a wound compared to mice (m) without the locus. The invention describes a novel method of identifying a gene involved in enhanced wound healing by identifying DNA microsatellite markers which can distinguish healer mice from non-healer mice and identifying microsatellite markers which segregate with enhanced wound healing in progeny of the mice, where a chromosomal locus containing at least one enhanced wound healing gene is identified. A method of treating a wound in a mammal is also disclosed. The new methods are useful for treating wounds, especially central and peripheral nerve wound. The methods of the invention are useful for restoring function after nerve injury in a manmal. (M) is useful as a mammalian model of enhanced wound healing, useful for identifying genes and gene products involved in enhanced wound healing, and to provide methods for wound healing. AA218691-219048 represent murine SAGE tags from C57BL/6 and MRL mice which are used to illustrate the method of the
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Matches 10; Conservative
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949 TTAATGTATC 958

10 TTAATGTATY 1

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RESULT 1368 AAZ18749

13-FEB-1998; 26-AUG-1998; 28-SEP-1998;

12-FEB-1999;

WO9941364-A2

Mus sp.

19-AUG-1999

Heber-Katz E;

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The present invention describes a polynucleotide (I) of 956 base pairs (bp) given in the specification. The polynucleotide, especially the 5'-rCTTT TTCTTY, sequence (II), is useful for site specific recombination, and introducing and removing desired genes into mammalian cells. (II) is also useful for transgenic work and as a recombination expressed mammalian protein(s) that initiate a novel type of site-specific recombination in mammalian cells. (I) represents a DNA fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Assay of genetic sequences based on triplex formation from double stranded analyte - and hybrid of anchor and reporter sequences, with reporter released if triplex formation occurs, used e.g. to identify bacteria.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Triple helix forming nucleotides 997-1007 of 23S rRNA gene.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           12.9%; Score 9.4; DB 1; Length 11; 90.9%; Pred. No. 1.2e+03; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                New polynucleotides isolated from the hamster Sp5 clone.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Seguence 11 BP; 0 A; 2 C; 0 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                     (GENO-) GENOTOX TESTING & CONSULTING HB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (PROF-) PROFILE DIAGNOSTIC SCI INC.
                                                                                                                                                                                                                                                                                                                                                             Claim 3; Page 14; 23pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAX14920 standard; DNA; 11 BP.
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                                                                                                   99WO-SE000573
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nes 10; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              From a hamster Sp5 clone
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                                                                                                                                                                                                                                Jensen D, Helleday T;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Wang C;
                                                                                                                                                                                                                                                                       WPI; 1999-620423/53.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4PI; 1999-130384/11.
                  W09953048-A1
                                                                                                   08-APR-1999;
                                                                                                                                              08-AFR-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               29-OCT-1992;
                                                          21-OCT-1999.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hepburn AG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAX14920;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This invention describes a novel non-WRL healer mouse (M) having at least one quantitative trait locus selected from those given in the specification, exhibiting an enhanced healing response to a wound compared to mice (M) without the locus. The invention describes a novel method of identifying a gene involved in enhanced wound healing by identifying a gene involved in enhanced wound healing by contributing microsatellite markers which segregate with enhanced wound healing in progeny of the mice, where a chromosomal locus containing at least one enhanced wound healing gene is identified. A method of treating a wound; a mammal is also disclosed. The new methods are useful for treating wounds, especially central and peripheral nerve wound. The methods of the invention are useful for restoring function after nerve injury in a mammal. (M) is useful as a mammalian model of enhanced wound healing, useful for identifying genes and gene products involved in enhanced wound healing, and to provide methods for wound healing are used to illustrate the method of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              mammalian model for enhanced wound healing - useful for identifying
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                    Wound healing; non-MRL healer mouse; quantitative trait locus; QTL; healing response; microsatellite marker; treatment; central nerve; peripheral nerve; nerve injury; SAGE tag; murine; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Chinese hamster; Sp5; mutant; site specific genetic recombination;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 13; Page 73; 136pp; English.
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98US-0097937P.
98US-0102051P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     enhanced wound healing genes
Murine MRL SAGE tag 1568982.
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nes 10; Conservative
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13-JAN-2000

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AAZ32222

RESULT 1370

Query Match

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                                The present sequence represents a potential triple-helix forming region. It can be used to demonstrate the assay of the invention. The assay comprises adding a sample containing double-stranded DNA test sequences, e.g. containing the present sequence, to an aqueous medium containing at least one complex of anchor DNA, attached to a solid support, and reporter DNA, where either a part of the anchor DNA or reporter DNA is designed to form a triple-strand structure with part of the test sequence. Triplex formation results in displacement of the reporter DNA which is detected as an indication of the presence of the DNA test sequence. The method is used to detect DNA sequences, particularly for identification of bacteria (by detecting genes for ribosomal RNA) in clinical samples, but also detection of oncogenes and Hepatitis B virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The present invention relates to a recombinant plasmid comprising a region with a nuclectide sequence capable of specifically binding to a sequence-specific DNA-binding molecule, a region with a nuclectide sequence capable of binding to a restriction enzyme and a restriction site for a restriction enzyme. The invention is useful for detecting the presence of initiation of transcription activity by RNA polymerase and for detecting the presence of sequence-specific DNA binding molecules
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Novel recombinant plasmid useful for determining the activity of DNA binding protein, and for detecting the activity of RNA polymerases in initiating transcription.
                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                                  12.9%; Score 9.4; DB 1; Length 11; 90.9%; Pred. No. 1.2e+03; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                          Sequence 11 BP; 0 A; 4 C; 0 G; 7 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA-binding; RNA polymerase; transcription; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                DNA binding protein recognition sequence #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Page 20-24; 98pp; English.
           Disclosure; Col 23-24; 168pp; English.
                                                                                                                                                                                                                                                                                                                                                                                     BP.
                                                                                                                                                                                                                                                                                                                                                                 99US-00344300
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                                                                                                                                                                                                                                                                                           918 TCTTTGCCTTT 928
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The present sequence is provided in a specification relating to an isolated subgenomic polynuclectide comprising a percoxisome proliferator—activator receptor (PpAR) delta binding element and an RXR binding element. The polynuclectide is useful for identifying potential therapeutic agents for cancer treatment and for ameliorating negative side effects of non-stroidal anti-inflammatory diseases (NSAIDS). Test compounds which increase transcription of PPARdelta protein, PPARdelta protein binding to a PPARdelta binding element, or expression of a reporter gene which is under the control of a PPARdelta binding element, are identified. These are candidates for use in encouraging cell are identified. These are candidates for use in a disease state such as allowed, anyotrophic lateral actions, or other muscle wasting diseases, autoimmune diseases, heart attack, stroke, ischaemic heart disease, kidney failure, septic shock, or a disease, in which the cell is infected with a pathogen, such as a virus, bacterium, fungus, mycoplasma, or protoace negative side effects of NSAIDs, stomach or intestines, or to ameliorate negative side effects of NSAIDs,
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                                                                                                                                                                                                                                                                                                  Human; peroxisome proliferator-activator receptor delta; PPARdelta; RXR; cytostatic; nootropic; neuroprotective; anti-HIV; cardiant; cerebroprotective; vasotropic; antiulcer; immunosuppressive; nophrotropic; antibacterial; antiviral; antifungal; protozoacide; non-steroidal anti-inflammatory disease; NSAID; infection; Alzheimer's disease; ALDS; muscle wasting disease; autoimmune disease; binding element; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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                                                                                                                                                                                                                                                 Human RXR binding element, SEQ ID NO: 28.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Vogelstein B;
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ВÞ.
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AAF75228 standard; DNA; 11
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                                                                                                                                                                09-MAY-2001
                                                                           AAF75228;
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C) le

12.9%; Score 9.4; DB 1; Length 11; 90.9%; Pred. No. 1.2e+03;

Query Match Best Local Similarity

Hofmann K;

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The invention relates to identifying (M1) genes in vitro that, in humans or animals, are important for skin ageing and/or skin stress by serial analysis of gene expression between mixtures of transcribed and optionally translated, genetically encoded factors (A) obtained from young and aged skin, to identify that genes that show strong differential useful for: identifying markers of skin ageing and/or stress; determining skin ageing and/or stress; and identifying or determining the effects of pharmaceutical not comentic agents for control of skin ageing. The present sequence is one of a group of human skin ageing/stress related expressed sequence tags (ABQ86246-ABQ87680) of the invention
                                                                                                                                                                                                                                            Identifying genes involved in skin stress and aging, useful e.g. in screening for cosmetic or therapeutic agents, based on differential gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12.9%; Score 9.4; DB 1; Length 11; 90.9%; Pred. No. 1.2e+03; cive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 11 BP; 0 A; 3 C; 1 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                       Claim 8; Page 82; 325pp; German.
                                                                                                 20-DEC-2001; 2001WO-EP015178.
                                                                                                                           03-JAN-2001; 2001DE-01000121
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Matches 10; Conservative
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                                        WO200253773-A2
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             Homo sapiens.
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                                                                                                                                                                                      Petersohn D,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The invention relates to identifying (M1) genes in vitro that, in humans or animals, are important for skin ageing and/or skin stress by serial analysis of gene expression between mixtures of transcribed and optionally translated, genetically encoded factors (A) obtained from young and aged skin, to identify that genes that show strong differential useful for: identifying markers of skin ageing and/or stress; determining skin ageing and/or stress; and identifying or determining the effects of pharmacoutical or cosmetic agents for control of skin ageing. The present sequence is one of a group of human skin ageing/stress related expressed sequence tags (ABQ86246-ABQ87680) of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Identifying genes involved in skin stress and aging, useful e.g. in screening for cosmetic or therapeutic agents, based on differential gene expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                    Human; skin ageing; skin stress; EST; expressed sequence tag; ss.
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                                                                                                                                                                                      Human skin stress/ageing related EST SEQ ID NO 74.
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                                                                                                   ABQ86319 standard; cDNA; 11 BP
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Matches 10; Conservative
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                 1 CCTGGTCAATT
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                                                                                                                                                                                                                                                                                                                                                                                                                           Petersohn D,
                                                                                                                                                                                                                                                   Homo sapiens.
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                                                                                                                              ABQ86319
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ID ABO8

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Human; skin ageing; skin stress; EST; expressed sequence tag; ss.
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                                                                                                                              ABQ86887 standard; cDNA; 11 BP.
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                                                                                                                        The invention relates to identifying (MI) genes in vitro that, in humans or animals, are important for skin ageing and/or skin stress by serial analysis of gene expression between mixtures of transcribed and optionally translated, genetically encoded factors (A) obtained from expression. (A) comprises protein or mRNAMs or their fragments. (MI) is useful for: identifying markers of skin ageing and/or stress; and identifying or determining the effects of skin ageing and/or stress; and identifying or determining the effects of shammaceutical or cosmetic agents for control of skin ageing. The present sequence is one of a group of human skin ageing/stress related expressed sequence tags (ABQ86246-ABQ87680) of the invention
                Identifying genes involved in skin stress and aging, useful e.g. in screening for cosmetic or therapeutic agents, based on differential gene
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                                                                                                                                                                                                                                                                                                                                                                                          Score 9.4; DB 1;
Pred. No. 1.2e+03;
0; Mismatches 1;
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                                                                                          Claim 8; Page 63; 325pp; German.
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Best Local Similarity 90.9%;
Matches 10; Conservative
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expression,
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ABQ87291/
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sequence is one of a group of human skin ageing/stress related expressed sequence tags (ABQ86246-ABQ87680) of the invention
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                                                                           Score 9.4; DB 1; Length 11 Pred. No. 1.2e+03; 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                  Human skin stress/ageing related EST SEQ ID NO 790.
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                                                Seguence 11 BP; 7 A; 1 C; 1 G; 2 T; 0 U; 0 Other;
                                                                                                            0; Mismatches
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                                                                              12.9%;
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                                                                                                                                                                                                                                                        ABQ87035 standard; cDNA; 11
                                                                                                                                                                                                                                                                                                                    10-SEP-2002 (first entry)
                                                                           Query Match
Best Local Similarity 90.9
Matches 10; Conservative
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Best Local Similarity
Matches 10; Conserv
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ABV65020/c
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The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression. (M1) is useful for identifying genes involved in skin homeostasis; to promotes skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitis; sunburn; psoriaais, scleroderma; ichthyosis; atopic dermatitis; acne; seborrhea; lupus erythematosus; rosaces; melanoma; basal cell carcinoma; and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag (EST) of the invention
                                                                                                                                                                                                                                                                                                                   In vitro identification of skin-expressed genes, useful for determining homeostasis and identifying cosmetic or pharmaceutical agents against
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12.9%; Score 9.4; DB 1; Length 11; 90.9%; Pred. No. 1.2e+03; cive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 11 BP; 8 A; 0 C; 2 G; 1 T; 0 U; 0 Other;
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                                                                                                     20-DEC-2001; 2001WO-EP015179.
                                                                                                                                              03-JAN-2001; 2001DE-01000127.
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Best Local Similarity
Matches 10; Conserv
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                                                             11-JUL-2002
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                                                                                                                                                                Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              In vitro identification of skin-expressed genes, useful for determining homeostasis and identifying cosmetic or pharmaceutical agents against
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ABV65020 standard; cDNA; 11 BP
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ABV66616 standard; cDNA; 11 BP.
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90.9%;
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Matches 10; Conservative
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                                                                                                                          Human skin EST 2806
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Human, skin, dermatological, vulnerary, antipsoriatic, antiseborrhaeic, immunosuppressive, antiinflammatory, cytostatic, SAGE, neurodermatitis, psoriasis, dermatitis, skin cancer, EST, expressed sequence tag, ss.
ABV69524 standard; cDNA; 11 BP.
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ABV66616

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Homo sapiens
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                                                                                                                                                The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression ($AGE) so as to identify skin-expressed genes and quantify their expression ($AGE) (M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin ichthyosis, specifically neurodermatitis; sunburn; psoriasis; scleroderma; ichthyosis; atopic dermatitis; acnes; seborrhea; lupus expressed screen of the can be used for the can be used to be used to can be used 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.
                                   In vitro identification of skin-expressed genes, useful for determining homeostasis and identifying cosmetic or pharmaceutical agents against
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Pred. No. 1.2e+03;
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                                                                                                                 Disclosure; Page 229; 1345pp; German
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90.9%;
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Best Local Similarity 90.9
Matches 10; Conservative
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WPI; 2002-590638/63
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                                                                              e.g. skin cancer.
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determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitis; sunburn; psoriasis; scleroderma; ichthyosis; atopic dermatitis; acne; seborrhea; lupus erythematosus; rosacea; melanoma; basal cell carcinoma; and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag (ESI) of the invention
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Best Local Similarity
...rhes 10; Conserve
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Human skin EST 720.
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immunosuppressive, antiinflammatory, cytostatic, SAGE, neurodermatitis,
psoriasis, dermatitis, skin cancer, EST, expressed sequence tag, ss.
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                                                                                           ABV68041 standard; cDNA; 11
                                                                                                                                       21-OCT-2002 (first entry)
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Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic;
immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis;
psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.
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ABV62934 standard; cDNA; 11

RESULT 1385

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The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression.

(M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitis; sumburn; psoriaris; scleroderma; ichthyosis; atopic dermatitis; sumburn; psoriaris; scleroderma; rosacea; melanoma; basal cell carcinoma; and carcinoma or sarcoma of the
                                                          The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression.

(M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitis; sunburn; psoriasis; scleroderma; ichthyosis; atopic dermatitis; acne; seborrhea; lupus erythematosus; rosacea; melanoma; basal cell carcinoma; and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag (BST) of the invention
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                          Claim 24; Page 260; 1345pp; German.
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Matches 10; Conservative
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                                                                                                                                                    Hofmann K;
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20-DEC-2001; 2001WO-EP015179
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Best Local Similarity 90.9
Matches 10; Conservative
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(first entry)

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Human, skin, dermatological, vulnerary, antipsoriatic, antiseborrhaeic, immunosuppressive, antiinflammatory, cytostatic, SAGE, neurodermatitis, psoriasis, dermatitis, skin cancer, EST, expressed sequence tag, ss.
                                     ABV65581 standard; cDNA; 11 BP
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                                     G; 1 T; 0 U; 0 Other;
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90.9%;
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90.9%;
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Best Local Similarity 
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Conservative
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Matches 10; Conservative
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skin. The present seque (EST) of the invention
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03-JAN-2001; 2001DE-01000127.
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Matches 10;
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Human, skin, dermatological, vulnerary, antipsoriatic, antiseborrhaeic, immunosuppressive, antiinflammatory, cytostatic, SAGE, neurodermatitis,

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Gaps

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                                                                                                                                                                                     In vitro identification of skin-expressed genes, useful for determining homeostasis and identifying cosmetic or pharmaceutical agents against e.g. skin cancer.
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psoriasis; dermatitis; skin cancer; EST; expressed sequence tag;
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Best Local Similarity
Matches 10; Conserv
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immunosuppressive, antiinflammatory, cytostatic, SAGE, neurodermatitis,
psoriasis, dermatitis, skin cancer, EST, expressed sequence tag, ss.
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20-DEC-2001; 2001WO-EP015179 Σ Conradt (HENK) HENKEL KGAA Petersohn D, 11-JUL-2002 RESULT 1394 ABV62773

03-JAN-2001; 2001DE-01000127

Hofmann K;

WPI; 2002-590638/63.

In vitro identification of skin-expressed genes, useful for determining homeostasis and identifying cosmetic or pharmaceutical agents against e.g. skin cancer.

Disclosure; Page 41; 1345pp; German.

The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression. (M1) is useful for identifying genes involved in skin homeostasis, to promotes skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin identifically neurodermatitis; sunburn; psoriasis, scleroderma; ichthyosis; atopic dermatitis; acne; seborrhea; lupus erythematosus; ichthyosis; melanoma; basal cell carcinoma; and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag

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ö The invention relates to in vitro identification (W1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression. (M1) is useful for identifying genes involved in skin homeostasis; to promotes skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitis; sunburn, psoriaeis, scleroderma; ichthyosis; atopic dermatitis; acne; seborrhea; lupus erythematosus; sosacea, melanoma; basal cell carcinoma; and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag Human, skin, dermatological, vulnerary, antipsoriatic, antiseborrhaeic, immunosuppressive, antiinflammatory, cytostatic, SAGE, neurodermatitis, psoriasis, dermatitis, skin cancer, EST, expressed sequence tag, ss. In vitro identification of skin-expressed genes, useful for determining homeostasis and identifying cosmetic or pharmaceutical agents against Gaps ; Score 9.4; DB 1; Length 11; Pred. No. 1.2e+03; 0; Mismatches 1; Indels Score 9.4; DB 1; Length 11; Pred. No. 1.2e+03; Sequence 11 BP; 1 A; 3 C; 0 G; 7 T; 0 U; 0 Other; Disclosure; Page 210; 1345pp; German. Hofmann K; BP. 20-DEC-2001; 2001WO-EP015179. 03-JAN-2001; 2001DE-01000127. 12.9%; 12.9%; 90.9%; ABV68868 standard; cDNA; 11 (first entry) Query Match Best Local Similarity 90.9 Matches 10; Conservative 915 Conradt M, ; 905 TCATTTTCTTT TCTTTTTTT Human skin EST 6654. WPI; 2002-590638/63. (HENK) HENKEL KGAA Local Similarity e.g. skin cancer. WO200253774-A2. Homo sapiens Petersohn D, 21-OCT-2002 11-JUL-2002. ABV68868; Query Match RESULT 1395 Best Loca Matches 8

ABV63019;

Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss. ABV66065 standard; cDNA; 11 BP. (first entry) 911 TCTTTGGTCTT 921 1 retriecter 11 Human skin EST 3851. WO200253774-A2 Homo sapiens 21-OCT-2002 RESULT 1397 ð 엄

Matches 10; Conservative

Query Match Best Local Similarity

The invention relates to in vitro identification (MI) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (\$AGE) so as to identify skin-expressed genes and quantify their expression. (MI) is useful for identifying genes involved in skin homeostasis, to promotes skin homeostasis and to the tagant (A) that maintains or promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitis; sunburn; psoriasis; scleroderma; inchthyosis; atophy dermatitis, anne, seborrhea; lupus exprhematosus; rosaces, melanoma; basal cell carcinoma; and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag Human, skin, dermatological, vulnerary, antipsoriatic, antiseborrhaeic, immunosuppressive, antiinflammatory, cytostatic, SAGE, neurodermatitis, psoriasis, dermatitis, skin cancer, EST, expressed sequence tag, ss. In vitro identification of skin-expressed genes, useful for determining homeostasis and identifying cosmetic or pharmaceutical agents against Sequence 11 BP; 0 A; 3 C; 1 G; 7 T; 0 U; 0 Other; X Disclosure; Page 47; 1345pp; German. Hofmann 20-DEC-2001; 2001WO-EP015179. 03-JAN-2001; 2001DE-01000127 ΣÌ of the invention Petersohn D, Conradt (HENK) HENKEL KGAA WPI; 2002-590638/63 Human skin EST 805 e.g. skin cancer. WO200253774-A2 Homo sapiens. 11-JUL-2002

12.9%; Score 9.4; DB 1; Length 11; 90.9%; Pred. No. 1.2e+03; ive 0; Mismatches 1; Indels BP. ABV70955 standard; cDNA; 11 (first entry) 0.9 Query Match Best Local Similarity 90.9 Matches 10, Conservative 920 TITGCCTTTTA 930 Trescritia 11 Human skin EST 8741 21-OCT-2002 ABV70955; RESULT 1398
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The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression.

(M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitis; sunburn; psoriasis; scleroderma; ichthyosis; atopic dermatitis; acne; seborrhea; lupus erythematosus; ichthyosis; melanoma; basal cell carcinoma; and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag In vitro identification of skin-expressed genes, useful for determining homeostasis and identifying cosmetic or pharmaceutical agents against e.g. skin cancer. Disclosure; Page 131; 1345pp; German Hofmann K; 20-DEC-2001; 2001WO-EP015179 03-JAN-2001; 2001DE-01000127 Conradt M, WPI; 2002-590638/63 (HENK) HENKEL KGAA Petersohn D, 11-JUL-2002

Human, skin, dermatological, vulnerary, antipsoriatic, antiseborrhaeic, immunosuppressive, antiinflammatory, cytostatic, SAGE, neurodermatitis, psoriasis, dermatitis, skin cancer, EST, expressed sequence tag, ss.

Homo sapiens

WO200253774-A2.

11-JUL-2002.

20-DEC-2001; 2001WO-EP015179

03-JAN-2001; 2001DE-01000127

(HENK) HENKEL KGAA

Hofmann K; Σ Petersohn D, Conradt

WPI; 2002-590638/63

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The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression (M1) is useful for identifying genes involved in skin homeostasis, to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitis; subburn; psoriasis; scleroderma;
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disorders, specifically neurodermatitis; sunburn; psoriasis; scleroderma; ichthyosis, atopic dermatitis; acne; seborornea; lupus erythematosus; rosacea; melanoma; basal cell carcinoma; and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag
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Best Local Similarity 90.9%;
Matches 10; Conservative
                                                                                                             (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 932 CCCCCCCTCTTC 942
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                                                                                                                                                                                                                                                                                                                                             Petersohn D, Conradt
                                                                                                                                    Human skin EST 5284
                                                                                                                                                                                                                                                                                                                      (HENK ) HENKEL KGAA
                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-590638/63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human skin EST 7980
                                                                                                                                                                                                                                                                                                                                                                                                                e.g. skin cancer.
                                                                                                                                                                                                                             WO200253774-A2.
                                                                                                                                                                                                        Homo sapiens
                                                                                                              21-OCT-2002
                                                                                                                                                                                                                                                   11-JUL-2002,
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                                                                                        ABV67498;
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                                           RESULT 1401
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ID ABV7
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DE Huma
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression. (M1) is useful for identifying genes involved in skin homeostasis; to promotes skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis and to test agent (A) that maintains or ichthyosis; atopic dermatitis; sunburn; psoriasis, scleroderma; lichthyosis; atopic dermatitis; acne; seborrhea; lupus erythematcsus; rosacea; melanoma; basal cell carcinoma; and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag (EST) of the invention
Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    In vitro identification of skin-expressed genes, useful for determining homeostasis and identifying cosmetic or pharmaceutical agents against
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 11 BP; 0 A; 2 C; 0 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hofmann K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 24; Page 254; 1345pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RXR binding site from clone X9TOP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ABV78654 standard; DNA; 11 BP
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مور مع عير م

Park BH,

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Novel CYPJAS polynucleotide useful for diagnosis and treatment of cancer, cardiovascular diseases, diabetes and AIDS, and for identifying
                                                                                                       The invention relates to sequence determination oligonucleotides for detecting polymorphic sites in a 5' flanking region of cytochrome P450 different sene. CYP2D6 enzymes are involved in the metabolism of many different xenobiotics. Human CYP2D6 gene is located on chromosome 22. The logonucleotides may be used as in situ hybridisation probes, in ligase-based sequenced determination, as components of diagnostic assays, as probes in sequence determination methods based on mismatches, as hybridisation-based diagnostic assays, and as components of diagnostic ability to metabolise certain drugs. The present sequence is a sense oligonucleotide used for detecting of human CYP2D6 gene 5' flanking region single nucleotide polymorphism (SNP)
hybridization probes, as components of diagnostic assays, or in ligase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human, CYP3AS, polymorphism; cancer; cardiovascular disease, diabetes; AIDS, African American; forensic marker; pharmacological; cytostatic; antidiabetic; anti-HIV; gene therapy; ds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The present invention relates to a new CYP3A5 polynucleotide encoding polypeptide, where the polynucleotide is capable of hybridising to a
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 9.4; DB 1; Length 11; Pred. No. 1.2e+03; 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human CYP3A5 gene polymorphic variant DNA sequence #21.
                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 11 BP; 1 A; 1 C; 3 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (EPID-) EPIDAUROS BIOTECHNOLOGIE AG
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                                                                   Claim 2; Page 23; 63pp; English
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     28-DEC-2000; 2000US-0258684P.
29-DEC-2000; 2000US-0258952P.
16-JAN-2001; 2001EP-00100172.
16-JAN-2001; 2001US-026899P.
16-AUG-2001; 2001EP-00118884.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        12.9%;
90.9%;
                          based sequence determination
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Best Local Similarity 90.9
Matches 10; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           polymorphisms.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              11-JUL-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 1405
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                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to a novel homozygous peroxisome proliferatoractivated receptor delta (PPARdelta) gene-defective cell line. The compositions of the invention have nootropic, neuroprotective, anti-HIV, cardiant, cytostatic, anti-HIAMMALOTY, immunosuppressive, and cerebroprotective activity. The cell lines may have a use in gene therapy. The methods and compositions are useful for treating inflammation and cancer and other disorders with increased cell proliferation or in which cells are dying prematurely such as Alzheimer's disease, AIDS, muscular dystrophy, autoimmune diseases, heart attack and stroke, improving fecundity and/or ameliorating toxic effects of nonstenoidal anti-inflammatory drugs. The sequence represents a PCR product of an oligonucleotide template that bound a fusion protein containing the DNA binding domain of RXR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human; cytochrome P450 2D6; CYP2D6; enzyme; detection; xenobiotic;
ligase-based sequenced determination; drug metabolism; chromosome 22; ss.
                                                                                                                                                                                                                           Homozygous PPAR gene-defective cell line, useful for treating inflammation and cancer and disorders associated with premature cell death such as Alzheimer's disease, AIDS, muscular dystrophy, autoimmune diseases and heart attacks.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human CYP2D6 gene polymorphic site 942 detecting sense 5' oligo.
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                                                                                                                                         ä
                                                                                                                                           Vogelstein
                                                                                                                                                                                                                                                                                                                                              Example 2; Fig 6; 33pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       27-AUG-2001; 2001WO-IB001544.
    27-FEB-2002; 2002WO-US003408
                                                27-FEB-2001; 2001US-0271412P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            12.9%;
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                                                                                             (UYJO ) UNIV JOHNS HOPKINS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAD34267 standard; DNA; 11
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Best Local Similarity 90.9
Matches 10; Conservative
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                                                                                                                                         Kinzler KW,
                                                                                                                                                                                    WPI; 2002-691649/74.
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07-MAR-2002

AAD34267;

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Gaps ö S

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12.9%; Score 9.4; DB 1; Length 12;
Best Local Similarity 90.9%; Pred. No. 1.2e+03;
Matches 10; Conservative 0; Mismatches 1; Indels Sequence 12 BP; 7 A; 0 C; 5 G; 0 T; 0 U; 0 Other;

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Gaps

931 TCCCTCCTCT 941

TICCICCICI 12 g à

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AAX14829

BP. AAX14829 standard; DNA; 12

AAX14829;

(first entry)

24-MAR-1999

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Gaps

Triple helix third strand of 23S rRNA gene nucleotides 212-223.

Triplex formation; DNA detection; triple helix; identification; bacteria;

oncogene; virus;

Synthetic. Escherichia coli.

US5861244-A.

19-JAN-1999

93US-00173489,

92US-00968436. 29-OCT-1992; (PROF-) PROFILE DIAGNOSTIC SCI INC.

Hepburn AG,

WPI; 1999-130384/11.

Assay of genetic sequences based on triplex formation from double stranded analyte - and hybrid of anchor and reporter sequences, with reporter released if triplex formation occurs, used e.g. to identify bacteria.

Disclosure; Col 21-22; 168pp; English.

The present sequence represents a polymucleotide that is able to form a triple helix with a double stranded sequence. Cytosine bases in the present can be replaced with 5-methylotosine for increased triplex tability. The present sequence is used in the assay of the invention, where it can be part of the anchor DNA or reporter DNA sequence. The assay comprises adding a sample containing double-stranded DNA test. DNA, attached to an aqueous medium containing at least one complex of anchor DNA, attached to a solid support, and reporter DNA, where either a part of the anchor DNA or reporter DNA, sets of complex of anchor DNA, attached to a solid support, and reporter DNA, where either a part of the anchor DNA or reporter DNA is designed to form a triple-strand structure with part of the test sequence. Triplex formation results in displacement of the reporter DNA which is detected as an indication of the presence of the DNA test sequence. The method is used to detect DNA sequences, particularly for identification of bacteria (by detecting genes for ribosomal RNA) in clinical samples, but also detecting on occosens and Hepatitis B virus

Sequence 12 BP; 0 A; 4 C; 0 G; 8 T; 0 U; 0 Other;

6 12.9%; Score 9.4; DB 1; Length 12; 90.9%; Pred. No. 1.2e+03; tive 0; Mismatches 1; Indels Local Similarity 90.5 Best Loca Matches

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Gaps

905 TCATTITCTT 915 2 TCCTTTTCTTT

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CYP3A5 gene. The invention is useful in an in vitro method for identifying a polymorphism. The invention is also useful for useful for diagnosing a disorder related to the presence of a molecular variant of a CYP3A5 or susceptibility to such a disorder, where the disorder is cancer, or diseases including cardiovascular diseases, diabetes and AIDS. The invention can further be used for the preparation of a diagnostic composition for diagnosing a disease in a subject having a genome comprising a variant allele of the CYP3A5 gene, where the subject is an African American. The molecules of the invention are as forensic markers and in pharmacological studies. The present nucleic acid sequence represents a human CYP3A5 gene polymorphism variant DNA sequence, as ; 0 Score 9.4; DB 1; Length 11; Pred. No. 1.2e+03; 0; Mismatches 1; Indels Sequence 11 BP; 1 A; 2 C; 1 G; 7 T; 0 U; 0 Other; Query Match Best Local Similarity 90.9%; Matches 10; Conservative 911 TCTTTGGTCTT 921

8×30000000000×8

1 TCTTTGATCTT 11 셤

1406

AAX14622 standard; DNA; 12 BP.

AAX14622,

(first entry) 24-MAR-1999 Triple helix forming nucleotides 6650-6661 of the c-myc gene.

Triple-helix forming region, Triplex formation, DNA detection, identification, bacteria, oncogene, virus, ds.

Homo sapiens

US5861244-A

19-JAN-1999

93US-00173489 22-DEC-1993; 92US-00968436 29-OCT-1992; (PROF-) PROFILE DIAGNOSTIC SCI INC.

Wang C; Hepburn AG,

WPI; 1999-130384/11.

Assay of genetic sequences based on triplex formation from double stranded analyte - and hybrid of anchor and reporter sequences, with reporter released if triplex formation occurs, used e.g. to identify bacteria.

Disclosure, Col 13-14; 168pp; English.

The present sequence represents a potential triple-helix forming region. It can be used to demonstrate the assay of the invention. The assay comprises adding a sample containing double-erranded DNA test sequences, e.g. containing the present sequence, to an aqueous medium containing at least one complex of anchor DNA, attached to a solid support, and reporter DNA, where either a part of the anchor DNA or reporter DNA is designed to form a triple-strand structure with part of the test sequence. Triplex formation results in displacement of the reporter DNA which is detected as an indication of the presence of the DNA test sequence. The method is used to detect DNA sequences, particularly for identification of bacteria (by detecting genes for ribosomal RNA) in clinical samples, but also detection of oncogenes and Hepatitis B virus AAX14622/
AAX14622/
AAX14622/
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AAX1 XXX
AAX1 24-M
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AXX 12-M
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Page 638

Mon Oct 18 14:40:13 2004

schultz1-899.rng

(first entry)

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                                                                                                                                                                                                                                                                                                                                                                                                                                 Identifying target genes in a microorganism (e.g. Escherichia coli) as a basis for anti-infective treatment comprises selecting potential targets known to be present and obtaining complementary (antisense) peptide nucleic acid sequences.
                                                                                                                                                                                                /*tag= a
/mod_base= OTHER
/note= "linked to AAB99988 by 8-amino-3,6-dioxaoctanoic
acid"
                                                                                                                              Peptide nucleic acid, PNA, antimicrobial, antibiotic, cationic peptide;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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                                                                                                         Antibacterial peptide nucleic acid oligonucleotide #57.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 12 BP; 3 A; 4 C; 2 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                        Wissenbach M;
                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 3; Page 35; 57pp; English.
                                          AAH23548 standard; DNA; 12 BP.
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                                                                                                                                                                                                                                                                                                                                  99DK-00001468.
99US-0159683P.
                                                                                                                                          antisense; disinfectant; ss.
                                                                                    (first entry)
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                                                                                                                                                                                                                                                                                                                                                                (PANT-) PANTHECO AS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
                                                                                                                                                                                                                                                                 WO200127262-A1
                                                                                                                                                                                      Key
modified_base
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                                                                                    03-AUG-2001
                                                                                                                                                                                                                                                                                      19-APR-2001
                                                                                                                                                               Synthetic.
                                                               AAH23548;
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ABI17707 standard; DNA; 12 BP.

RESULT 1409 ABI17707/c ID ABI17707 XX AC ABI17707

ABI17707

954 GTATCGCTACC 964 [||||| ||||| GTATCACTACC 11

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                                                  SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Oligonucleotide primer SEQ ID NO 317680 for detecting SNP TSC0028168.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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nes 10; Conservative
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99889, ABH00010-ABH9988 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic form at from WIPO at
                                                                                                                                                                                                                              our or oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                                                                                                                                                                                                                                                  Claim 1; SEQ ID NO 324104; 29pp + Sequence Listing; German.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         was obtained in electronic format from Wi
ftp.wipo.int/pub/published_pct_sequences
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                                        06-APR-2001; 2001WO-IB000713.
                                                                                2000DE-01019173
                                                                                                                                                                  Olek A, Piepenbrock C,
                                                                                                                        (EPIG-) EPIGENOMICS AG
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                                                                                07-APR-2000;
18-OCT-2001
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/ Match 12.9%; Score 9.4; DB 1; Length 12; Local Similarity 90.9%; Pred. No. 1.2e+03; nes 10; Conservative 0; Mismatches 1; Indels
                                                                                             905 TCATTTTCTT 915
    Query Match
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ABH74276 standard; DNA; 12 (first entry) 22-FEB-2002 ABH74276;

BP.

Oligonucleotide primer SEQ ID NO 274261 for detecting SNP TSC0003493.

SNP, single nucleotide polymorphism, human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Homo sapiens

WO200177384-A2.

18-OCT-2001

06-APR-2001; 2001WO-IB000713

07-APR-2000; 2000DE-01019173

(EPIG-) EPIGENOMICS

Olek A,

WPI; 2001-657177/75

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Berlin

. 13 Set of oligonucleotides, useful for diagnosis and cell typing, 1.1

designed to detect single-nucleotide polymorphisms and cytosine methylation status.

Claim 1; SEQ ID NO 274261; 29pp + Sequence Listing; German.

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and oytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABE99989, ABF00010-ABE99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences

Sequence 12 BP; 3 A; 5 C; 0 G; 4 T; 0 U; 0 Other;

Gaps ö / Match 12.9%; Score 9.4; DB 1; Length 12; Local Similarity 90.9%; Pred. No. 1.2e+03; nes 10; Conservative 0; Mismatches 1; Indels Query Match Best Loca Matches

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RESULT 1412

ABI25548 standard; DNA; 12 BP.

(first entry) 22-FEB-2002 ABI25548;

Oligonucleotide primer SEQ ID NO 325521 for detecting SNP TSC0032582

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Homo sapiens

WO200177384-A2

18-OCT-2001

06-APR-2001; 2001WO-IB000713.

07-APR-2000; 2000DE-01019173

(EPIG-) EPIGENOMICS AG

Berlin K; Olek A, Piepenbrock C,

WPI; 2001-657177/75

Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine methylation status.

Claim 1; SEQ ID NO 325521; 29pp + Sequence Listing; German

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073

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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                           Oligonucleotide primer SEQ ID NO 302598 for detecting SNP TSC0020076.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 1; SEQ ID NO 302598; 29pp + Sequence Listing; German.
                                                                                ABI02625 standard; DNA; 12 BP.
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                                                                                                                                              (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                     SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                               Gaps
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                                                                                                           12.9%; Score 9.4; DB 1; Length 12; 90.9%; Pred. No. 1.2e+03; ive 0; Mismatches 1; Indels
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                                                                              Sequence 12 BP; 2 A; 0 C; 2 G; 8 T; 0 U; 0 Other;
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                                               ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                                                          ABI27706 standard; DNA; 12
                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                                                                                           Query Match 12.9
Best Local Similarity 90.9
Matches 10, Conservative
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF99989, ABF00010-ABF99989, ABF00010-ABF99989, and ABI00010-ABF82073 data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 12 BP; 6 A; 3 C; 0 G; 3 T; 0 U; 0 Other;
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                              Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                                                                                                                                                                          Claim 1; SEQ ID NO 279208; 29pp + Sequence Listing; German.
                                                                                                                                                                       Berlin K;
                                                                                                                            07-APR-2000; 2000DE-01019173
                                                                                                         06-APR-2001; 2001WO-IB000713
                                                                                                                                                                       Olek A, Piepenbrock C,
                                                                                                                                                                                            WPI; 2001-657177/75.
                                                                                                                                                  (EPIG-) EPIGENOMICS
                                                               WO200177384-A2
                                            Homo sapiens,
                                                                                     18-OCT-2001
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genemic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but fur was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences

Gaps 0 Query Match 12.9%; Score 9.4; DB 1; Length 12; Best Local Similarity 90.9%; Pred. No. 1.2e+03; Matches 10; Conservative 0; Mismatches 1; Indels Sequence 12 BP; 8 A; 4 C; 0 G; 0 T; 0 U; 0 Other;

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913 TTTGGTCTTTG 923 12 TTTGGTTTTG

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ABI06994 standard; DNA; 12 BP. 22-FEB-2002 (first entry) ABI06994; RESULT 1416 ABI06994

Oligonucleotide primer SEQ ID NO 306967 for detecting SNP TSC0022272.

SNP; single nuclectide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Homo sapiens

WO200177384-A2

18-OCT-2001

06-APR-2001; 2001WO-IB000713

07-APR-2000; 2000DE-01019173

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF99999, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic formmat from WFPO at fire wipo.int/pub/published_pot_sequences
                                                                                                                                           Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                                                                                                                                                          Claim 1; SEQ ID NO 306967; 29pp + Sequence Listing; German.
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                                                             Berlin
                                                             Piepenbrock C,
                  (EPIG-) EPIGENOMICS AG.
                                                                                                        WPI; 2001-657177/75.
                                                             olek A,
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Query Match 12.9%; Score 9.4; DB 1; Length 12; Best Local Similarity 90.9%; Pred. No. 1.2e+03; Matches 10; Conservative 0; Mismatches 1; Indels 924 CCTTTTATCCC 934

2 CCTTTTATTCC 12

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Gaps

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BP. RESULT 1417

Oligonucleotide primer SEQ ID NO 307293 for detecting SNP TSC0022421.

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

06-APR-2001; 2001WO-IB000713.

07-APR-2000; 2000DE-01019173.

Berlin

Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.

Claim 1; SEQ ID NO 307293; 29pp + Sequence Listing; German.

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Sequence 12 BP; 8 A; 0 C; 3 G; 1 T; 0 U; 0 Other;

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              acid (PNA) oligomers for detecting single nuclectide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABR09989, ABR00010-ABR9989, ABR00010-ABR9989, ABR00010-ABR9989, ABR00010-ABR99999, ABR00010-ABR99999 and ABI00010-ABR82073 at represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form par of the printed specification, but ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretraeted genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABE9989, ABF00010-ABE9989, ABF00010-ABE9989, ABF00010-ABE9989, and ABI00010-ABE3073 represent the oligomers described in the invention. NOTE: The sequence was obtained in electronic format from WIPO at the printed specification, but ftp.wipo.int/pub/published_pct_sequences
invention describes novel oligonucleotide primers or peptide nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                                   Score 9.4; DB 1; Length 12;
Pred. No. 1.2e+03;
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                                                                                                                                                                                                                                                                                                  SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                      Oligonucleotide primer SEQ ID NO 337518 for detecting SNP TSC0039907.
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RESULT 1420 ABI38144 ... ເນ

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABF99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                                                                     SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardicvascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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ABI38144 standard; DNA; 12 BP.
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                                                         Set of oligonucleotides, useful for diagnosis and cell typing, : designed to detect single-nucleotide polymorphisms and cytosine
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory,

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central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF0010-ABF99989, ABF0010-ABF99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at figure.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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designed to detect single-nucleotide polymorphisms and cytosine
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC09989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                          SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                 Oligonucleotide primer SEQ ID NO 357695 for detecting SNP TSC0050739.
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                                                                                                                                                                                                                                                                                                                                                                SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, contral nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF9989, ABH00010-ABH99999 and ABI00010-ABI82073 trepresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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12.9%; Score 9.4; DB 1; Length 12;
Best Local Similarity 90.9%; Pred. No. 1.2e+03;
Matches 10; Conservative 0; Mismatches 1; Indels
                             Claim 1; SEQ ID NO 377057; 29pp + Sequence Listing; German
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99899, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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             RESULT 1431
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                                                                                                                                                                                                                                                                                    SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                Oligonucleotide primer SEQ ID NO 366719 for detecting SNP TSC0055936.
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was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABF9989, and ABT00010-ABI32073 are represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from NPO at
                                                                                                                                                                                                                                                       SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                  Oligonuclectide primer SEQ ID NO 267339 for detecting SNP TSC0000119.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and oycosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99889, ABF00010-ABF99889, ABH00010-ABH99889 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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                                                            bet or oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                              Claim 1; SEQ ID NO 268821; 29pp + Sequence Listing; German.
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central nervous system; gastrointestinal; respiratory; immune; metabolic.
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designed to detect single-nucleotide polymorphisms and cytosine
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les 10; Conservative
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and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastroinfestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF999989, ABF00010-ABF999989, ABF00010-ABF999989, ABF00010-ABF999989, ABF00010-ABF999989, ABF00010-ABF99989, ABF00010-ABF9998999, ABF00010-ABF99989, ABF000010-ABF99989, ABF0000010-ABF99989, ABF000010-ABF99989, ABF0000010-ABF99989, ABF00000000-ABF99989, ABF999989, ABF999989, ABF999989, ABF999989, ABF999
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Pred. No. 1.2e+03;
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Matches 10; Conservative
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                            Oligonucleotide primer SEQ ID NO 322266 for detecting SNP TSC0030767.
                                      Gaps
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0; Mismatches 1; Indels
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12.9%;
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ABH73304/c
ID ABH73304 standard; DNA; 12 XX
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ABI22293/c
ID ABI22293
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ABH73304;

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This invention describes novel oligonucleotide primars or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABE99989, ABF00010-ABE99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                   SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                    Oligonucleotide primer SEQ ID NO 273289 for detecting SNP TSC0003130.
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Matches 10; Conservative
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99999, ABF00010-ABF99999, ABH00010-ABH99999 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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designed to detect single-nucleotide polymorphisms and cytosine
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WPI; 2001-657177/75.

WO200177384-A2

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                                                                                                                  This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, contral nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF9989, ABF00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                  Claim 1; SEQ ID NO 277207; 29pp + Sequence Listing; German.
                                                                                                                                                                                                                                                                                                                                                                                                                          12.9%; Score 9.4; DB 1; Length 12; 90.9%; Pred. No. 1.2e+03;
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Matches 10; Conservative
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-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at the printed specification, ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                    Gaps
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to detect single-nucleotide polymorphisms and cytosine
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                                                                                                                                             12.9%; Score 9.4; DB 1; Length 12; 90.9%; Pred. No. 1.2e+03; ive 0; Mismatches 1; Indels
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Best Local Similarity 90.9
Matches 10; Conservative
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ABI04185/c
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo int/pub/published_pct_sequences
                                                                                                                                                                                                                         SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                        Oligonucleotide primer SEQ ID NO 280456 for detecting SNP TSC0008655.
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Best Local Similarity 90.9
Matches 10; Conservative
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12 TIGGTIGAATG
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AB137376
ID AB137376
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           SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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.; 0 This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, contral nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99899, ABH00010-ABH99989 and ABI00010-ABI82073 tapeses the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but thp.wipo.int/pub/published_pct_sequences Gaps Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status. ; Claim 1; SEQ ID NO 292025; 29pp + Sequence Listing; German. 12.9%; Score 9.4; DB 1; Length 12; 90.9%; Pred. No. 1.2e+03; ive 0; Mismatches 1; Indels Sequence 12 BP; 4 A; 0 C; 5 G; 3 T; 0 U; 0 Other; Berlin K; 07-APR-2000; 2000DE-01019173 Query Match
Best Local Similarity 90.9
Matches 10; Conservative 930 ATCCCTCCTCT 940 Olek A, Piepenbrock C, (EPIG-) EPIGENOMICS AG 12 ATCCATCCTCT WPI; 2001-657177/75

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ABI46663 standard; DNA; 12 ABI46663; RESULT 1445 ABI46663

BP.

Oligonucleotide primer SEQ ID NO 346636 for detecting SNP TSC0007729. (first entry) 22-FEB-2002

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Homo sapiens

WO200177384-A2.

18-OCT-2001

06-APR-2001; 2001WO-IB000713

07-APR-2000; 2000DE-01019173

(EPIG-) EPIGENOMICS AG.

Berlin K; Olek A, Piepenbrock C,

WPI; 2001-657177/75.

Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.

Claim 1; SEQ ID NO 346636; 29pp + Sequence Listing; German.

Y.

. 0 This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 the represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from NIPO at SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic. Oligonucleotide primer SEQ ID NO 352851 for detecting SNP TSC0048131. Gaps Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine methylation status. . 0 Claim 1; SEQ ID NO 352851; 29pp + Sequence Listing; German. Length 12; 12.9%; Score 9.4; DB 1; Length 12 90.9%; Pred. No. 1.2e+03; ative 0; Mismatches 1; Indels Sequence 12 BP; 1 A; 0 C; 3 G; 8 T; 0 U; 0 Other; ftp.wipo.int/pub/published_pct_sequences Berlin 06-APR-2001; 2001WO-IB000713. 07-APR-2000; 2000DE-01019173. 878/c ABI52878 standard; DNA; 12 22-FEB-2002 (first entry) Matches 10; Conservative 913 TTTGGTCTTTG 923 Tricerrire 11 Olek A, Piepenbrock C, WPI; 2001-657177/75. (EPIG-) EPIGENOMICS Query Match Best Local Similarity WO200177384-A2. Homo sapiens 18-OCT-2001. ABI52878; ABIS2378/14
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF9989, ABH0010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence was obtained in electronic formmat from WIPD at the printed specification, but the wipo.int/pub/published_pct_sequences

(first entry)

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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                           Oligonucleotide primer SEQ ID NO 379882 for detecting SNP TSC0063508.
                             ABI79909 standard; DNA; 12 BP.
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RESULT 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Oligonucleotide primer SEQ ID NO 359157 for detecting SNP TSC0051484.
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                                                                             12.9%; Score 9.4; DB 1; Length 12; 90.9%; Pred. No. 1.2e+03; ive 0; Mismatches 1; Indels
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                                Sequence 12 BP; 6 A; 3 C; 0 G; 3 T; 0 U; 0 Other;
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Best Local Similarity 90.9
Matches 10; Conservative
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Berlin K;

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Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from MFPO at

Sequence 12 BP; 1 A; 3 C; 0 G; 8 T; 0 U; 0 Other;

Query Match

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Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine methylation status.

WPI; 2001-657177/75.

Claim 1; SEQ ID NO 295707; 29pp + Sequence Listing; German.

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99899 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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12.9%; Score 9.4; DB 1; Length 12;
Best Local Similarity 90.9%; Pred. No. 1.2e+03;
Matches 10; Conservative 0; Mismatches 1; Indels
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Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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12.9%; Score 9.4; DB 1; Length 12; 90.9%; Pred. No. 1.2e+03; tive 0; Mismatches 1; Indels
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                                               Local Similarity 90.5
les 10; Conservative
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ID ABH71841

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range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC099889, ABF00010-ABF09989, ABF00010-ABF99898, ABF00010-ABF99898, ABF00010-ABF99989, ABF00010-ABF99989, and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                         SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic diseorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99899, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic formmat from WIPO at
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, coingomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF9989, ABH00010-ABH9989 and ABI00010-ABF82073 terpresent the oligomers described in the invention. NOTE: The sequence date for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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designed to detect single-nucleotide polymorphisms and cytosine
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                                                                                       SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                             Oligonucleotide primer SEQ ID NO 304479 for detecting SNP TSC0020963.
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                                                                 This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligoners for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligoners are also used for detecting cell type differentiation. ABC00010-ABC99899, ABF0010-ABF99899, ABF0010-ABF99899 and ABI0010-ABF92031 represent the oligoners described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SNP; single nuclectide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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methylation status.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Pred. No. 1.2e+03;
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Sequence 12 BP; 2 A; 0 C; 4 G; 6 T; 0 U; 0 Other;
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les 10; Conservative
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peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                         Homo sapiens
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ABI40821/
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                                                                                                                                                       SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                               Oligonucleotide primer SEQ ID NO 332613 for detecting SNP TSC0037029.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0001 PABC99989, ABR00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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Pred. No. 1.2e+03;
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ftp.wipo.int/pub/published_pct_sequences
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06-APR-2001; 2001WO-IB000713.
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Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine
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                                  designed to detect methylation status.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABT00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from NIPO at
acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99389, ABC0010-ABE99389, ABC0010-ABE99389 and ABI0010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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Pred. No. 1.2e+03;
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to detect single-nucleotide polymorphisms and cytosine
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                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1; SEQ ID NO 340794; 29pp + Sequence Listing; German.
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                                                                                                      Berlin K;
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                        (EPIG-) EPIGENOMICS AG
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Sequence 12 BP; 2 A; 0 C; 2 G; 8 T; 0 U; 0 Other;

This invention describes novel oligonucleotide primers or peptide nucleic

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(first entry)

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acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory. Central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC09989, ABF00010-ABF99989, ABF00010-ABF99989, ABF00010-ABF99989 and ABI00010-ABF99989, represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo int/pub/published_pct_sequences
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                                                                                                                                                                                 SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                  Oligonucleotide primer SEQ ID NO 347644 for detecting SNP TSC0010372.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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              Score 9.4; DB 1; Length 12;
Pred. No. 1.2e+03;
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Score 9.4; DB 1; Length 12;
Pred. No. 1.2e+03;
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ABI47671 standard; DNA; 12 BP.

RESULT 1465 AB147671/c ID AB14767

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABF00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                  Set of oligonucleotides, useful for diagnosis and cell designed to detect single-nucleotide polymorphisms and methylation status.
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oligomers are also used for detecting cell type differentiation. ABC00010 -ABC9989, ABF0010-ABF99899, ABF0010-ABF99899, ABF0010-ABF99899, ABF0010-ABF99899, ABF0010-ABF99999, ABF0010-ABF99999, ABF0010-ABF99999, ABF0010-ABF9999, ABF99999, ABF9999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Oligonucleotide primer SEQ ID NO 372491 for detecting SNP TSC0059419.
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                                                                                                                                                                                                             Match 12,9%; Score 9.4; DB 1; Length 12; Local Similarity 90.9%; Pred. No. 1.2e+03; es 10; Conservative 0; Mismatches 1; Indels
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                                                                                                                                                                     Sequence 12 BP; 1 A; 0 C; 4 G; 7 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     was obtained in electronic format from W. ftp.wipo.int/pub/published_pct_sequences
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methylation status.
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                                                                                                                                                                                                                       Oligonucleotide primer SEQ ID NO 373230 for detecting SNP TSC0059917.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC09989, ABF00010-ABF9989, ABH00010-ABH99899 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; 8s; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Oligonucleotide primer SEQ ID NO 360034 for detecting SNP TSC0051895.
                                                                                                                                                                                                                                                                                                                                                        set or oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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Pred. No. 1.2e+03;
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, contral nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF9989, ABH00010-ABH9989 and ABI00010-ABI82073 represent the oligomers described in the invention. NoTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from MIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SNP; single nucleotide polymorphism; human; diagnosie; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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ftp.wipo.int/pub/published_pct_sequences
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                                                                            Piepenbrock C,
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06-APR-2001; 2001WO-IB000713.

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretracted genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99999, ABF00010-ABH99999 and ABI00010-ABI82073 are present the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretracted genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99899, ABH00010-ABH99989 and ABI00010-ABI82073 tepseent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Oligonucleotide primer SEQ ID NO 268286 for detecting SNP TSC0001040.
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Claim 1; SEQ ID NO 317619; 29pp + Sequence Listing; German.
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                                                                                                                                                                                                                                                                                                                                                 Score 9.4; DB 1; Length 12;
Pred. No. 1.2e+03;
0; Mismatches 1; Indels
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il Similarity 90.9%;
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                                                                                                                                                                                                                                                                                                                                                                                          SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                           Oligonucleotide primer SEQ ID NO 318631 for detecting SNP TSC0028776.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Set of oligonuclectides, useful for diagnosis and cell typing, i designed to detect single-nuclectide polymorphisms and cytosine methylation status.
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                                                                 Length 12;
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                                                                                                 1; Indels
                              Sequence 12 BP; 4 A; 0 C; 7 G; 1 T; 0 U; 0 Other;
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                                                             Score 9.4; DB 1;
Pred. No. 1.2e+03;
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ftp.wipo.int/pub/published_pct_sequences
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Best Local Similarity 90.9
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                                                                                                                                                                                                                                                                                                              SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                               Oligonucleotide primer SEQ ID NO 294777 for detecting SNP TSC0016278.
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Best Local Similarity
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Oligonucleotide primer SEQ ID NO 275486 for detecting SNP TSC0003907.

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABE99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABT00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                                                                                     Claim 1; SEQ ID NO 275486; 29pp + Sequence Listing; German.
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                                                                                                                                                                                                                                                                                                                        methylation status.
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                                                                                                                                 This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABE99989 and ABI00010-ABE82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form mart of the printed specification, but the wipo int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Oligonucleotide primer SEQ ID NO 303591 for detecting SNP TSC0020541.
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                                                    Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                            Claim 1; SEQ ID NO 276111; 29pp + Sequence Listing; German.
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 Piepenbrock C,
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oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at fire printed specification, but ftp.wipo.int/pub/published_pct_sequences
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ABI05670 standard; DNA; 12
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Best Local Similarity 90.9
Matches 10; Conservative
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretraeted genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99889, ABF00010-ABF99899, ABH00010-ABH99989 and ABI00010-ABI82073 tepresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic formmat from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                          SND; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Oligonucleotide primer SEQ ID NO 283371 for detecting SNP TSC0011278.
                                            Oligonucleotide primer SEQ ID NO 332438 for detecting SNP TSC0036911.
                                                                                                                                                                                                                                                                                                                                                                                                     uer or oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                                                                                                                                                                                SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                    Oligonucleotide primer SEQ ID NO 331775 for detecting SNP TSC0036472.
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Pred. No. 1.2e+03;
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ABI32465 standard; DNA; 12
                                                                                                                                                                      ABI31802 standard; DNA; 12
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   Best Local Similarity 90.9
Matches 10; Conservative
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                                                   943 ATTGGTTTAAT 953
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             designed to detect methylation status.
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ABI32465/C
ID ABI32465
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AC ABI32465
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Gaps

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designed to detect single-nucleotide polymorphisms and cytosine methylation status.

Claim 1; SEQ ID NO 284886; 29pp + Sequence Listing; German

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, cancer also used for addiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9999, ABF00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but they.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                     Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                                                                                                                                                                                                                                Claim 1; SEQ ID NO 283371; 29pp + Sequence Listing; German.
                                                                                                                                                          Berlin K;
                                       06-APR-2001; 2001WO-IB000713
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18-OCT-2001
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                                                         Query Match
12.9%; Score 9.4; DB 1; Length 12;
Best Local Similarity 90.9%; Pred. No. 1.2e+03;
Matches 10; Conservative 0; Mismatches 1; Indels
Sequence 12 BP; 5 A; 0 C; 6 G; 1 T; 0 U; 0 Other;
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929 TATCCCTCCTC 939 12 TATCTCTCTC 2 ò 셤

ABH84893 standard; DNA; 12 (first entry) 22-FEB-2002 ABH84893; 1484

BP.

Oligonucleotide primer SEQ ID NO 284886 for detecting SNP TSC0012041.

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Homo sapiens

WO200177384-A2.

18-OCT-2001

06-APR-2001; 2001WO-IB000713.

07-APR-2000; 2000DE-01019173.

(EPIG-) EPIGENOMICS AG

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Set of oligonucleotides, useful for diagnosis and cell typing, is

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at

Sequence 12 BP; 3 A; 0 C; 4 G; 5 T; 0 U; 0 Other;

Gaps ö 12.9%; Score 9.4; DB 1; Length 12; 90.9%; Pred. No. 1.2e+03; tive 0; Mismatches 1; Indels Query Match
Best Local Similarity 90.9
Matches 10, Conservative

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945 TGGTTTAATGT 955 2 regrraaarer 12 ð g

ABH85174 standard; DNA; 12 ABH85174;

BP.

22-FEB-2002 (first entry)

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic. Oligonucleotide primer SEQ ID NO 285167 for detecting SNP TSC0012179.

Homo sapiens

WO200177384-A2.

18-OCT-2001

06-APR-2001; 2001WO-IB000713

07-APR-2000; 2000DE-01019173

(EPIG-) EPIGENOMICS AG.

Berlin K; Olek A, Piepenbrock C,

WPI; 2001-657177/75.

Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine methylation status,

Claim 1; SEQ ID NO 285167; 29pp + Sequence Listing; German.

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABH82073 RESULT 1485
ABH85174
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                  Oligonucleotide primer SEQ ID NO 376689 for detecting SNP TSC0061933.
                                                                                                                                                                                                                                                                                                                       06-APR-2001; 2001WO-IB000713.
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                                                                    ABI76716 standard; DNA; 12
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                                                                                                                                                                                                                                                                                                                                                               SNP, single nucleotide polymorphism, human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                      Oligonucleotide primer SEQ ID NO 289523 for detecting SNP TSC0013972.
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Pred. No. 1.2e+03;
0; Mismatches 1; Indels
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                                                                     Sequence 12 BP; 0 A; 0 C; 4 G; 8 T; 0 U; 0 Other
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Matches 10; Conservative
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Piepenbrock C,
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Best Local Similarity
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Oligonucleotide primer SEQ ID NO 365377 for detecting SNP TSC0055076.

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Gaps

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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Best Local Similarity 90.9
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic formmat from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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designed to detect single-nucleotide polymorphisms and cytosine
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                                                                                                                                designed to detect methylation status.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genemic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                                                                                                                      SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                     Oligonucleotide primer SEQ ID NO 323078 for detecting SNP TSC0031211.
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12.9%; Score 9.4; DB 1; Length 12;
Best Local Similarity 90.9%; Pred. No. 1.2e+03;
Matches 10; Conservative 0; Mismatches 1; Indels
                               Score 9.4; DB 1; Length 12;
Pred. No. 1.2e+03;
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Seguence 12 BP; 1 A; 2 C; 0 G; 9 T; 0 U; 0 Other;
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RESULT 1493 ABI01560

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABB9989, ABF00010-ABB9989, ABF00010-ABB9989, ABF00010-ABB99999, ABF00010-ABB99999 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence was obtained in electronic format from WIPO at the printed specification, but the wipo.int/pub/published_pct_sequences

ABI01560;

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic formmat from WIPD at
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                                                                                                                                                                                                                                                                                                                                 Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form mart of the printed specification, but ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                      SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                              Oligonucleotide primer SEQ ID NO 301533 for detecting SNP TSC0019538.
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                                            Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine methylation status.
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central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABC99989, ABF00010-ABC9989, ABF000010-ABC9989, ABF000010-ABC9989, ABF000010-ABC9989, ABF000010-ABC9989, ABF000010-ABC9989, ABF000010-ABC998, ABF000000000-ABC
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Pred. No. 1.2e+03;
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This invention describes novel oligonucleotide primers or peptide nucleic acid (FNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory,

Claim 1; SEQ ID NO 330541; 29pp + Sequence Listing; German.

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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
              Oligonucleotide primer SEQ ID NO 341672 for detecting SNP TSC0042176.
                                                                                                                                                                                                                                                                                                                                    Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status,
                                                                                                                                                                                        06-APR-2001; 2001WO-IB000713.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99889 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                                                                                                                               SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                              ABI41696 standard; DNA; 12 BP.
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Berlin K;

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, contral nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 trepresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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Claim 1; SEQ ID NO 341672; 29pp + Sequence Listing; German.
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Best Local Similarity 90.9
Matches 10, Conservative
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ABI41699 standard; DNA; 12

RESULT 1499

(first entry)

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ABI41699;

AB141699/ ID AB14 XX AB14 XX AB14 DT 22-F

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06-APR-2001; 2001WO-IB000713.
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                                                                   This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretraeted genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                   Claim 1; SEQ ID NO 346807; 29pp + Sequence Listing; German.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 1502
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(first entry)

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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                             Oligonucleotide primer SEQ ID NO 355632 for detecting SNP TSC0049746.
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ABI55659 standard; DNA; 12 BP.
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                                                                       ABI55659
                 RESULT 1504
ABI55659/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but fip.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                  SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                                         Oligonucleotide primer SEQ ID NO 351950 for detecting SNP TSC0047593.
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                                                                  12.9%; Score 9.4; DB 1; Length 12; 90.9%; Pred. No. 1.2e+03; ive 0; Mismatches 1; Indels
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                                       Sequence 12 BP; 5 A; 2 C; 0 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   12.9%; Score 9.4; DB 1; 90.9%; Pred. No. 1.2e+03; ive 0; Mismatches 1;
was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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                                                                                 Local Similarity 90.9
es 10; Conservative
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Best Local Similarity 90.9
Matches 10; Conservative
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                         Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                                                                                                               Claim 1; SEQ ID NO 355632; 29pp + Sequence Listing; German.
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Berlin K;
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nuclectide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC09989, ABC0010-ABE9989, ABF00010-ABE99989, ABF00010-ABE99989, ABF00010-ABE99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence was obtained in electronic formmat from WFPO at the printed specification, but the wipo.int/pub/published_pct_sequences
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Best Local Similarity 90.9
Matches 10, Conservative
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                  Piepenbrock C,
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ABI80840
장. 월
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Pred. No. 1.2e+03;
0; Mismatches 1; Indels
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90.9%;
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Matches 10; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         methylation status.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP)
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and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABF00010-ABF999899, ABF00010-ABF99989, ABF00010-ABF99989, ABF00010-ABF99989, ABF00010-ABF99989, ABF00010-ABF99989, ABF00010-ABF99989, ABF00010-ABF99989, ABF00010-ABF99989, ABF00010-ABF99989, ABF00010-ABF9998999, ABF00010-ABF99989, ABF000010-ABF99989, ABF000010-ABF99989, ABF000010-ABF99989, ABF000010-ABF99989, ABF000010-ABF99989, ABF000010-ABF99989, ABF000010-ABF99989, ABF000010-ABF99998, ABF000010-ABF99989, ABF000010-ABF9999, ABF000010-ABF99999, ABF000010-ABF9999, ABF000010-ABF9999, ABF000010-ABF9999, ABF000010-ABF9999, ABF000010-ABF9999
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Score 9.4; DB 1; Length 12;
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                                                                                                                                                                                          SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                 Oligonucleotide primer SEQ ID NO 270470 for detecting SNP TSC0002151.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genemic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99899, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABH3073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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oligonucleotides, useful for diagnosis and cell typing, ied to detect single-nucleotide polymorphisms and cytosine
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                                                                                                                                               Claim 1; SEQ ID NO 325293; 29pp + Sequence Listing; German.
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-ABC99989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                Sequence 12 BP; 4 A; 1 C; 2 G; 5 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                  SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                         Oligonucleotide primer SEQ ID NO 308456 for detecting SNP TSC0023023.
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99889, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at
                                                                                                                                                                    Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                                                                                                                                                                            Claim 1; SEQ ID NO 314441; 29pp + Sequence Listing; German
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07-APR-2000; 2000DE-01019173
                                                                                 Olek A, Piepenbrock C,
                                       (EPIG-) EPIGENOMICS
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. Query Match 12.9%; Score 9.4; DB 1; Length 12; Best Local Similarity 90.9%; Pred. No. 1.2e+03; Matches 10; Conservative 0; Mismatches 1; Indels 908 TTTTCTTTGGT 918 ò Db

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Gaps

||||| ||||| 12 TTTTATTTGGT 2

ABI14588 standard; DNA; 12 ABI14588; RESULT 1518 ABI14588

BP.

Oligonucleotide primer SEQ ID NO 314561 for detecting SNP TSC0026429. (first entry) 22-FEB-2002

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Homo sapiens

WO200177384-A2.

18-OCT-2001

06-APR-2001; 2001WO-IB000713.

07-APR-2000; 2000DE-01019173

(EPIG-) EPIGENOMICS AG.

ĸ Berlin Olek A, Piepenbrock C,

WPI; 2001-657177/75.

Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.

Claim 1; SEQ ID NO 314561; 29pp + Sequence Listing; German.

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              This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99889, ABF00010-ABF99899, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic formmat from WIPO at
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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ABI41018 standard; DNA; 12
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Matches 10; Conservative
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                             Oligonucleotide primer SEQ ID NO 361123 for detecting SNP TSC0052454.
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designed to detect single-nucleotide polymorphisms and cytosine
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            ABI61150 standard; DNA; 12
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                                                                                                                                                                                                                                                                                                                    SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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designed to detect single-nucleotide polymorphisms and cytosine
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                                      12.9%; Score 9.4; DB 1; Length 12; 90.9%; Pred. No. 1.2e+03; ive 0; Mismatches 1; Indels
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            BP; 7 A; 0 C; 3 G; 2 T; 0 U; 0 Other;
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                                             This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, contral nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF99999, ABH00010-ABH99999 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic formmat from WIPO at
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Claim 1; SEQ ID NO 361123; 29pp + Sequence Listing; German.
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DDT 22-F
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KW PEPP KW
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KW Centl
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919 CTTTGCCTTTT 929

11 Crrrrccrrrr 1

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WPI; 2001-657177/75
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF99989, ABH00010-ABF99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic formmat from WIPO at fitte.wipo.int/pub/published_pot_sequences
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Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine methylation status.
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range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC00989, ABC0010-ABE9989, ABM00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form par of the printed specification, but was obtained in electronic format from WIPO at fire printed specification, but firm without the printed specification, but form with the printed specification of the printed specification.
                                                                                                                                                                                                                                                                                                                                                                                                                                        SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                                                                                                                                                             SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; 8s; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF0010-ABF99989, ABH0010-ABH99989 and ABI00010-ABI82073 targressent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from NIPO at
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                                                                                      SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                             Oligonucleotide primer SEQ ID NO 325328 for detecting SNP TSC0032508.
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Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine

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                                                                                              This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF9989, ABH00010-ABH99999 and ABI00010-ABI82073 tepresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Oligonucleotide primer SEQ ID NO 278193 for detecting SNP ISC0005779.
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                                                                                                                                                                                   12.9%; Score 9.4; DB 1; Length 12; 90.9%; Pred. No. 1.2e+03; ive 0; Mismatches 1; Indels
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ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                            peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Oligonucleotide primer SEQ ID NO 333884 for detecting SNP TSC0037810.
                                                                                                           Oligonucleotide primer SEQ ID NO 329309 for detecting SNP TSC0034878
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acid (FNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and range of diseases including immune system, gastrointestinal, respiratory. Central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99889, ABC0010-ABC99889 and ABI0010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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peptide nucleic acid, cytosine methylation; cardiovascular, primer; ss;
central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            oet or oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                                                                                                                                                                                                   WPI; 2001-657177/75.
                                 designed to detect s
methylation status.
                  WPI; 2001-657177/75
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretraeted genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascilar and metabolic disconders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABC0010-ABF9989, ABH00010-ABH9989 and ABI00010-ABI82073 tepresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically precreated genomic DNA. The oligomucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic discorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC09989, ABC0010-ABC99898, ABC0010-ABC99898, ABC0010-ABC099898 and ABI0010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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igned to detect single-nucleotide polymorphisms and cytosine
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12.9%; Score 9.4; DB 1; Length 12;
Best Local Similarity 90.9%; Pred. No. 1.2e+03;
Matches 10; Conservative 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             was obtained in electronic format from Wi
ftp.wipo.int/pub/published_pct_sequences
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                       (EPIG-) EPIGENOMICS AG
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Sequence 12 BP; 8 A; 0 C; 4 G; 0 T; 0 U; 0 Other;

This invention describes novel oligonuclectide primers or peptide nucleic

(first entry)

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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                    Oligonucleotide primer SEQ ID NO 287999 for detecting SNP TSC0013339.
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                                                                                                                                                                                                                                                                                                         (EPIG-) EPIGENOMICS AG
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              ABH88006;
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                                                                                                                                                                                                                                                                                        SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                          Gaps
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         Score 9.4; DB 1; Length 12;
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0; Mismatches 1; Indels
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         Query Match
Best Local Similarity
Matches 10; Conserv
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Berlin K;

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acid (PNA) oligomers for detecting single nuclectide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99889, ABC0010-ABC99898 and ABL00010-ABC809873 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo.int/pub/published_pct_sequences
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ABH88006 standard; DNA; 12 BP.

RESULT 1538 ABH88006/c ID ABH88006

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Matches
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                                                                                                                                                                                                 This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99899, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from wIPO at
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Oligonucleotide primer SEQ ID NO 343536 for detecting SNP TSC0010582.
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                                                                                                                                         oligonuclectides, useful for diagnosis and cell typing, :ed to detect single-nucleotide polymorphisms and cytosine
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                                         06-APR-2001; 2001WO-IB000713.
                                                           07-APR-2000; 2000DE-01019173.
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Best Local Similarity 90.5
Matches 10; Conservative
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                                                                                                    Piepenbrock C,
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                                                                                                                                                                methylation status.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The
                                                                                                                                                                                                                               This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and oytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99889, ABF00010-ABF99989, ABH0010-ABF99989, and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic formmat from WIPO at ftp.wipo.int/pub/published_pct_sequences
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Set of oligonucleotides, useful for diagnosis and cell typing, i
designed to detect single-nucleotide polymorphisms and cytosine
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                                                                                                                                                                   Claim 1; SEQ ID NO 343536; 29pp + Sequence Listing; German.
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Pred. No. 1.2e+03;
0; Mismatches 1; Indels
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                                               designed to detect Emethylation status.
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oligomers are also used for detecting cell type differentiation, ABC000100-ABC99989, ABF0010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at fixed specification, but ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Pred. No. 1.2e+03;
0; Mismatches 1; Indels
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12.9%; Score 9.4; DB 1; Length 12;
Best Local Similarity 90.9%; Pred. No. 1.2e+03;
Matches 10; Conservative 0; Mismatches 1; Indels
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, contral nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo.int/pub/published_pct_sequences
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ABI59468 standard; DNA; 12
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930 ATCCCTCTCT 940
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                  1 ATCCCTTCTCT 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                      SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Oligonucleotide primer SEQ ID NO 373728 for detecting SNP TSC0060292.
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                                                                                                                                                                                                                                                                      designed to detect a methylation status.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically prerreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC09989, ABF00010-ABF09989, ABF00010-ABF9989, ABF00010-ABF9989, ABF00010-ABF99989 and ABI00010-ABF82073 data for this patent did not form part of the printed specification, but typ.wipo.int/pub/published_pct_sequences
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                                             This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABR00010-ABH99989 and ABI00010-ABH99989 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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Claim 1; SEQ ID NO 379572; 29pp + Sequence Listing; German.
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Sequence 12 BP; 7 A; 3 C; 1 G; 1 T; 0 U; 0 Other;
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                                                                                                                                                                         SNP; single nuclectide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                         Oligonucleotide primer SEQ ID NO 305619 for detecting SNP TSC0021533.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF9989, ABH00010-ABH9989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent din ot form part of the printed specification, but was obtained in electronic format from WIPO at
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Best Local Similarity 90.9%; Pred. No. 1.2e+03;
Matches 10; Conservative 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                       methylation status.
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                 Homo sapiens.
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
   oligonucleotides are used for diagnosis and/or prognosis of cancer and a
                                                                                                                                                                                                                                                                                                                                                                                                                                               Oligonuclectide primer SEQ ID NO 285363 for detecting SNP TSC0012260.
                                                                                                                                                                               12.9%; Score 9.4; DB 1; Length 12; 90.9%; Pred. No. 1.2e+03; ive 0; Mismatches 1; Indels
                                                                                                                                                   Sequence 12 BP; 4 A; 1 C; 4 G; 3 T; 0 U; 0 Other;
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Best Local Similarity 90.3%,
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                                                                                                                                              This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genemic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABH32073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Oligonucleotide primer SEQ ID NO 309422 for detecting SNP TSC0023520.
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                                                          oligonucleotides, useful for diagnosis and cell typing, is to detect single-nucleotide polymorphisms and cytosine
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                                                                                                                    Claim 1; SEQ ID NO 333159; 29pp + Sequence Listing; German.
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Best Local Similarity 90.9
Matches 10; Conservative
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Olek A, Piepenbrock C,
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                                                          Set of oligonucleotidesigned to detect methylation status.
                             WPI; 2001-657177/75
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, oardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABE99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 tepresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                                                                                                                                               Claim 1; SEQ ID NO 285363; 29pp + Sequence Listing; German.
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ftp.wipo.int/pub/published_pct_sequences
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W0200177384-A2
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ABH88274/c
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                                                                                                                                                                                                      SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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            Gaps
                                                                                                                                                                                   Oligonucleotide primer SEQ ID NO 310465 for detecting SNP TSC0023992.
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Pred. No. 1.2e+03;
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90.9%;
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Best Local Similarity 90.9
Matches 10; Conservative
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Best Local Similarity
Matches 10; Conserv
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ABI11455 standard; DNA; 12 BP.

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                                                  SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Oligonucleotide primer SEQ ID NO 311428 for detecting SNP TSC0024493
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Pred. No. 1.2e+03;
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Best Local Similarity 90.9
Matches 10; Conservative
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                                                                                                              Olek A, Piepenbrock C,
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and merabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                                Claim 1; SEQ ID NO 313488; 29pp + Sequence Listing; German
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                        Gaps
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                                                                                              Sequence 12 BP; 5 A; 0 C; 2 G; 5 T; 0 U; 0 Other;
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Matches 10; Conservative
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                                                                                                                                                             SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                               Oligonucleotide primer SEQ ID NO 350874 for detecting SNP TSC0046946.
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                               ABIS0901 standard; DNA; 12 BP.
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RESULT 1560
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12.9%; Score 9.4; DB 1; Length 12; 90.9%; Pred. No. 1.2e+03; tive 0; Mismatches 1; Indels

Local Similarity 90.9

Best Loca Matches

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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99999, ABF00010-ABF99999, ABH00010-ABH99999 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cycosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF0010-ABF9989, ABH0010-ABH9989 and ABI0010-ABI82073 trepresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from MIPO at
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                         Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                                                              Claim 1; SEQ ID NO 356516; 29pp + Sequence Listing; German.
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ftp.wipo.int/pub/published_pct_sequences
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Sequence 12 BP; 2 A; 3 C; 0 G; 7 T; 0 U; 0 Other;

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 tepresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Oligonucleotide primer SEQ ID NO 381959 for detecting SNP TSC0004779.
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                                                                                                                                                                                                                                                                                                                                SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                 Oligonucleotide primer SEQ ID NO 268523 for detecting SNP TSC0001198.
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12.9%; Score 9.4; DB 1; Length 12; 90.9%; Pred. No. 1.2e+03;
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RESULT 1566 ABH72306/c

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acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABE09989, ABE00010-ABE9989, ABE00010-ABE9989, ABE00010-ABE9989, ABE00010-ABE9989, and on the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at the printed specification, but ftp.wipo.int/pub/published_pct_sequence
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                                                                                                                                                                                                                                                                   SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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designed to detect single-nucleotide polymorphisms and cytosine
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Pred. No. 1.2e+03;
0; Mismatches 1; Indels
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ABH72306 standard; DNA; 12
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Best Local Similarity 90.9
Matches 10; Conservative
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99899, ABH00010-ABH99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Oligonucleotide primer SEQ ID NO 335135 for detecting SNP TSC0038619.
                                tides, useful for diagnosis and cell typing, i
single-nucleotide polymorphisms and cytosine
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                                                                                                         Claim 1; SEQ ID NO 329412; 29pp + Sequence Listing; German.
                                                                                                                                                                                                                                                                                                                                                                                                             Score 9.4; DB 1; Length 12;
Pred. No. 1.2e+03;
0; Mismatches 1; Indels
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Matches 10; Conservative
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                                      oligonucleotides,
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WPI; 2001-657177/75
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central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99899, ABF00010-ABF99899 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at fig. wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                       Gaps
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                                                                                                                                                                               Score 9.4; DB 1; Length 12;
Pred. No. 1.2e+03;
0; Mismatches 1; Indels
                                                                                                                                                Sequence 12 BP; 1 A; 1 C; 0 G; 10 T; 0 U; 0 Other;
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Best Local Similarity 90.9
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Query Match
Best Local Similarity 90.9%; Pred. No. 1.2e+03;
Matches 10; Conservative 0; Mismatches 1; Indels

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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99889, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic formmat from WIPD at
                                             SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
               Oligonucleotide primer SEQ ID NO 353351 for detecting SNP TSC0048475.
                                                                                                                                                                                                                                                                                                                                                                                                        Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                                                                                                                                                         SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                           Oligonucleotide primer SEQ ID NO 344909 for detecting SNP TSC0043756.
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    12.9%; Score 9.4; DB 1; Length 12; 90.9%; Pred. No. 1.2e+03; tive 0; Mismatches 1; Indels
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ABI53378 standard; DNA; 12

RESULT 1572

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ABI53378;

ABI53378/ ID ABI5 XX AC ABI5 XX DT 22-F

06-APR-2001; 2001WO-IB000713. 07-APR-2000; 2000DE-01019173.

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, contral nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                              Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                                                                                                               Claim 1; SEQ ID NO 360645; 29pp + Sequence Listing;
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                                                                 (EPIG-) EPIGENOMICS
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABF99989 and ABL00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but

Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.

Berlin K;

Olek A, Piepenbrock C,

WPI; 2001-657177/75.

(EPIG-) EPIGENOMICS AG

07-APR-2000; 2000DE-01019173

Claim 1; SEQ ID NO 365497; 29pp + Sequence Listing; German.

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretraeted genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99899, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 tepresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic formmat from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SNP, single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Oligonucleotide primer SEQ ID NO 365497 for detecting SNP TSC0055166.
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           Claim 1; SEQ ID NO 375327; 29pp + Sequence Listing; German.
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                                                                                                                                                                                                                                  Sequence 12 BP; 7 A; 2 C; 0 G; 3 T; 0 U; 0 Other;
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                            Oligonucleotide primer SEQ ID NO 292806 for detecting SNP TSC0015368.
                                           ABH92813 standard; DNA; 12
                                                                                                22-FEB-2002
                                                                    ABH92813;
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                                                                                                                                                                                                                                                                                                                                   SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                 Score 9.4; DB 1; Length 12;
Pred. No. 1.2e+03;
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Best Local Similarity 90.9%; Pred. No. 1.2e+03;
Matches 10; Conservative 0; Mismatches 1; Indels
                                        Sequence 12 BP; 7 A; 3 C; 0 G; 2 T; 0 U; 0 Other;
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was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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Best Local Similarity 90.9%;
Matches 10; Conservative 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    designed to detect methylation status.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 the september of this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; 8s; central nervous system; gastrointestinal; respiratory; immune; metabolic.
central nervous system; gastrointestinal; respiratory; immune; metabolic.
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0; Mismatches 1; Indels
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                                                                                                                                                                                                              methylation status.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 was obtained in electronic format from W1 ftp.wipo.int/pub/published_pct_sequences
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and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99899, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but thp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and merabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABF99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                     Sequence 12 BP; 3 A; 0 C; 2 G; 7 T; 0 U; 0 Other;
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC09989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI32073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at
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                                                                                                                                  This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP)
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set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                            Claim 1; SEQ ID NO 333767; 29pp + Sequence Listing; German.
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-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, contral nervous system, cardiovascular and metabolic disorders. The coligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF9989, ABH0010-ABH99989 and ABI0010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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designed to detect single-nucleotide polymorphisms and cytosine
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         SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Homo sapiens

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, contral nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Pred. No. 1.2e+03;
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                                                             Olek A, Piepenbrock C,
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ö This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nuclectide polymorphisms (SNP) and cytosine methylation status in chemically prereated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99889, ABF00010-ABF99899, ABF00010-ABF99898 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo.int/pub/published_pct_sequences SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic. Gaps Oligonucleotide primer SEQ ID NO 373985 for detecting SNP TSC0060439. . 0 Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status. Claim 1; SEQ ID NO 373985; 29pp + Sequence Listing; German. Query Match
12.9%; Score 9.4; DB 1; Length 12;
Best Local Similarity 90.9%; Pred. No. 1.2e+03;
Matches 10; Conservative 0; Mismatches 1; Indels

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Claim 1; SEQ ID NO 358914; 29pp + Sequence Listing; German.

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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                               Oligonucleotide primer SEQ ID NO 362507 for detecting SNP TSC0006608.
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designed to detect single-nucleotide polymorphisms and cytosine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      claim 1; SEQ ID NO 362507; 29pp + Sequence Listing; German.
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            ABI62534 standard; DNA; 12
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                                                                                                                                                                                                                                                                                                                         SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                              Oligonucleotide primer SEQ ID NO 375661 for detecting SNP TSC0061372.
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                                        Score 9.4; DB 1; Length 12;
Pred. No. 1.2e+03;
0; Mismatches 1; Indels
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               Sequence 12 BP; 8 A; 4 C; 0 G; 0 T; 0 U; 0 Other;
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                                            Query Match
Best Local Similarity 90.9
Matches 10; Conservative
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Matches 10; Conservative
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Berlin K;

Piepenbrock C,

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The cligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99999, ABF00010-ABF99999, ABH00010-ABH99999 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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943 ATTGGTTTAAT 953

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                       set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                                       Claim 1; SEQ ID NO 364547; 29pp + Sequence Listing; German.
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Best Local Similarity 90.9
Matches 10; Conservative
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range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99389, ABF00010-ABF99989, ABF00010-ABF99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form par of the printed specification, but was obtained in electronic format from WIPO at fire wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF9989, ABH00010-ABF99989 and ABI00010-ABF8073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but
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Best Local Similarity 90.9
Matches 10; Conservative
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12.9%; Score 9.4; DB 1; Length 12; 90.9%; Pred. No. 1.2e+03;

Query Match Best Local Similarity

Seguence 12 BP; 1 A; 1 C; 3 G; 7 T; 0 U; 0 Other;

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                                                                                                                                                                                                                                                       SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                        Oligonucleotide primer SEQ ID NO 268667 for detecting SNP TSC0001288.
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designed to detect single-nucleotide polymorphisms and cytosine
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AC ABI18684;
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                                                      Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                            Oligonucleotide primer SEQ ID NO 318657 for detecting SNP TSC0028791.
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Pred. No. 1.2e+03;
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                                                        This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF99899, ABF00010-ABF99899, ABF00010-ABF99899, ABF00010-ABF99899 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                              Claim 1; SEQ ID NO 269622; 29pp + Sequence Listing; German.
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                                                                                                                                                                                                                                                                          Sequence 12 BP; 2 A; 0 C; 3 G; 7 T; 0 U; 0 Other;
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90.9%;
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data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABEC99899, ABF00010-ABE99899, ABH00010-ABE99899 and ABI00010-ABE82073 represent the oligomers described in the invention, NOTE: The sequence

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Homo sapiens.
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                                                                                                                               SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                         Oligonucleotide primer SEQ ID NO 284714 for detecting SNP ISC0011956.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99999, ABF00010-ABF99999, ABH00010-ABH99999 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at fire wipo.int/pub/published_pct_sequences
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peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                              Piepenbrock C,
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   (EPIG-) EPIGENOMICS AG
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                                                             WPI; 2001-657177/75
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This invention describes novel oligonucleotide primers or peptide nucleic

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acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and oytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and range of diseases including immune system, gastrointestinal, respiratory central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABB9989, ABF00010-ABB9989, ABF00010-ABB99999 and ABI0010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                                                          12.9%; Score 9.4; DB 1; Length 12; 90.9%; Pred. No. 1.2e+03;
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                                                                                                                                                                                                                                                                                                                                                                          0; Mismatches
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, contral nervous system, asriovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABE99989, ABF00010-ABE99989, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                  SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                          Oligonuclectide primer SEQ ID NO 313978 for, detecting SNP TSC0026057
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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Best Local 10; Conservative
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                     ABI14005;
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                                                                                                                                                                                                                                                                                                                                                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                 Oligonucleotide primer SEQ ID NO 288708 for detecting SNP TSC0013638.
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                  Length 12;
                  Query Match
12.9%; Score 9.4; DB 1; Length 12
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Matches 10; Conservative 0; Mismatches 1; Indels
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ABI14005 standard; DNA; 12
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Matches 10; Conservative
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, contral nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABH99989 and ABI00010-ABI82073 trepresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                          oligonucleotides, useful for diagnosis and cell typing, ied to detect single-nucleotide polymorphisms and cytosine
                                                                                                                                                                                                                                                 Claim 1; SEQ ID NO 342960; 29pp + Sequence Listing; German.
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Best Local Similarity 90.9
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ABI56583 standard; DNA; 12 BP.

RESULT 1614 ABI56583 22-FEB-2002 (first entry)

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ABI56583;

Sequence 12 BP; 0 A; 5 C; 0 G; 7 T; 0 U; 0 Other;

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99889, ABF00010-ABF99899, ABH00010-ABH99989 and ABI00010-ABF32073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic formmat from WIPO at

Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine methylation status.

Claim 1; SEQ ID NO 352749; 29pp + Sequence Listing; German.

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The
                                                                   SND; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
Oligonucleotide primer SEQ ID NO 356556 for detecting SNP TSC0050181.
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Berlin K;

Piepenbrock C,

olek A,

WPI; 2001-657177/75

(EPIG-) EPIGENOMICS AG

06-APR-2001; 2001WO-IB000713 07-APR-2000; 2000DE-01019173

WO200177384-A2.

18-OCT-2001.

Homo sapiens

Oligonucleotide primer SEQ ID NO 352749 for detecting SNP TSC0048074.

(first entry)

22-FEB-2002

ABI52776;

946 GGTTTAATGTA 956

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oligomers are also used for detecting cell type differentiation. ABC00010 -ABC39989, ABF0010-ABF99889, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                            Sequence 12 BP; 8 A; 2 C; 0 G; 2 T; 0 U; 0 Other;
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                                                                                                                                                   was obtained in electronic format from Wiftp.wipo.int/pub/published_pct_sequences
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, cointral nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invantion. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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Query Match

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Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine

designed to detect amethylation status.

Berlin K;

Piepenbrock C,

olek A,

WPI; 2001-657177/75

07-APR-2000; 2000DE-01019173.

SNP TSC0052120.

(EPIG-) EPIGENOMICS AG

Claim 1; SEQ ID NO 267588; 29pp + Sequence Listing; German.

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99889, ABF00010-ABF99899, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                          SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Oligonucleotide primer SEQ ID NO 360538 for detecting
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC09989, ABF00010-ABF09989, ABF00010-ABF99989, ABF00010-ABF99989, ABF00010-ABF99989 and ABI00010-ABF9073 represent the oligomers described in the invention. NOTE: The sequence was obtained in electronic formmat from WIPO at the printed specification, but the wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                     This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligoners for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immine system, gastrointestinal, respiratory, oligoners are also used for detecting cell type differentiation. ABC00010-ABC9989, ABR00010-ABF9989, ABR00010-ABF9989, ABR00010-ABF9989, ABR00010-ABF9999, ABR00010-ABF99999, ABR00010-ABF99999, ABR00010-ABF99999, ABR00010-ABF99999, ABR00010-ABF99999, ABR00010-ABF99999, ABR00
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Oligonucleotide primer SEQ ID NO 268403 for detecting SNP TSC0001101.
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Claim 1; SEQ ID NO 267619; 29pp + Sequence Listing, German.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 12.9%; Score 9.4; DB 1; Length 12; Best Local Similarity 90.9%; Pred. No. 1.2e+03; Matches 10; Conservative 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                            ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                              Oligonucleotide primer SEQ ID NO 320867 for detecting SNP TSC0029940.
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Score 9.4; DB 1; Length 12;
Pred. No. 1.2e+03;
0; Mismatches 1; Indels
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12.9%;
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               Local Similarity 90.3
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Best Local Similarity 90.9
Matches 10; Conservative
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Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.

Berlin K;

Olek A, Piepenbrock C,

WPI; 2001-657177/75

(EPIG-) EPIGENOMICS AG

06-APR-2001; 2001WO-IB000713. 07-APR-2000; 2000DE-01019173

WO200177384-A2

18-OCT-2001.

Homo sapiens.

Claim 1; SEQ ID NO 306691; 29pp + Sequence Listing; German

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ABH74828/C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; Ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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ABI39499 standard; DNA; 12
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                                                                                                                                                                                                              This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nuclectide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but
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                                                                                   Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine
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                                                                                                                                                                                                                                                                                                                                                                                                                                               was obtained in electronic format from Wiftp.wipo.int/pub/published_pct_sequences
  Berlin K;
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Olek A, Piepenbrock C,
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PMA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretracted genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic discorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABF99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo.int/pub/published_pot_sequences Claim 1; SEQ ID NO 339472; 29pp + Sequence Listing; German. DB 1; Length 12; Sequence 12 BP; 3 A; 0 C; 5 G; 4 T; 0 U; 0 Other;

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC099889, ABC00010-ABE99889, ABH00010-ABE99899, ABH00010-ABE99989, and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence was obtained in electronic formmat from WIPO at the printed specification, but the wipo.int/pub/published_pct_sequences
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                                                                                          SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                   Oligonucleotide primer SEQ ID NO 316279 for detecting SNP TSC0027369.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC009989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073
                                                                                                                                                                                                                                           acid (PNA) oligomers for detecting single nuclectide polymorphisms (SNP) and oytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010 ABC99989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABH82073 the patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                             invention describes novel oligonucleotide primers or peptide nucleic
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designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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ABI44709 standard; DNA; 12 BP.
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Best Local Similarity 90.5
Matches 10; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at the print plant from the will be the print from the ftp.wipo.int/pub/published_pct_sequences
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12.9%; Score 9.4; DB 1; Length 12;
Best Local Similarity 90.9%; Pred. No. 1.2e+03;
Matches 10; Conservative 0; Mismatches 1; Indels
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Best Local Similarity 90.5.
Thes 10; Conservative
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99889, ABF00010-ABF99899, ABH00010-ABH99999 and ABI00010-ABI82073 capesent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic formmat from WIPO at the printed specification, but the wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                     SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                  Oligonucleotide primer SEQ ID NO 369646 for detecting SNP TSC0057765.
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Matches 10; Conservative
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                                                      ABI69673 standard;
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RESULT 1633
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99889, ABF00010-ABF99899, ABH00010-ABH99899 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo int/pub/published_pct_sequences
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Query Match 12.9%; Score 9.4; DB 1; Length 12; Best Local Similarity 90.9%; Pred. No. 1.2e+03; Matches 10; Conservative 0; Mismatches 1; Indels

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Sequence 12 BP; 6 A; 0 C; 3 G; 3 T; 0 U; 0 Other;

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99899, ABR00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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ABI22516 standard; DNA; 12
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                                                                                                                                                          SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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             ABH76419 standard; DNA; 12
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RESULT 1639

AB126786

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Homo sapiens
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                                                                                                                                                                         SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                               Oligonucleotide primer SEQ ID NO 326759 for detecting SNP TSC0033267.
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ABI26786 standard; DNA; 12
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, contral nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic formmat from WIPO at
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                            onucleotides, useful for diagnosis and cell typing, : detect single-nucleotide polymorphisms and cytosine
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                                  oligonucleotides, useful for diagnosis
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central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF0010-ABE9989 and ABI0010-ABE82073 represent the oligomers described in the invention. NoTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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Best Local Similarity 90.9%; Pred. No. 1.2e+03;
Matches 10; Conservative 0; Mismatches 1; Indels
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory,

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99889, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPD at
                                                                      SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                   Oligonucleotide primer SEQ ID NO 367308 for detecting SNP TSC0056273.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                    SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                                       Oligonucleotide primer SEQ ID NO 348089 for detecting SNP TSC0000612.
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ABI67335 standard; DNA; 12
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06-APR-2001; 2001WO-IB000713

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99899, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form par of the printed specification, but the was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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methylation status.
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Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.

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Piepenbrock C,

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WPI; 2001-657177/75.

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acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010 ABC99989, ABF00010-ABP9989, ABF00010-ABP9989
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                                                                                      This invention describes novel oligonucleotide primers or peptide nucleic
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                          Claim 1; SEQ ID NO 358728; 29pp + Sequence Listing; German
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12.9%; Score 9.4; DB 1; Length 12; 90.9%; Pred. No. 1.2e+03;
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                               Oligonucleotide primer SEQ ID NO 326038 for detecting SNP TSC0032865.
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                                          ABI26065 standard; DNA; 12
                                                                                                   22-FEB-2002 (first entry)
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                                                                                                                                                                                                                                                                                                                                                    SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                         Oligonuclectide primer SEQ ID NO 320853 for detecting SNP TSC0029932.
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                                                                    12.9%; Score 9.4; DB 1; Length 12; 90.9%; Pred. No. 1.2e+03; ive 0; Mismatches 1; Indels
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Pred. No. 1.2e+03;
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                                        Sequence 12 BP; 3 A; 0 C; 6 G; 3 T; 0 U; 0 Other;
was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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Best Local Similarity 90.9%;
Matches 10; Conservative
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Best Local Similarity 90.9
Matches 10; Conservative
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting call type differentiation. ABC0010-ABC9989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo.int/pub/published_pot_sequences
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC099889, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at
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                                                           set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                          Claim 1; SEQ ID NO 280028; 29pp + Sequence Listing; German.
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Pred. No. 1.2e+03;
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central nervous system; gastrointestinal; respiratory; immune; metabolic.
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and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABE99989, ABF0010-ABE99989, ABF0010-ABE99989, ABF0010-ABE99989, ABF0010-ABE99989 and ABI0010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Best Local Similarity 90.9%; Pred. No. 1.2e+03;
Matches 10; Conservative 0; Mismatches 1; Indels
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Conservative 0; Mismatches 1; Indels
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic discorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at first into int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Oligonucleotide primer SEQ ID NO 377538 for detecting SNP TSC0062375.
                                                                                                                                                                                                                                                                              Set of oligonucleotides, useful for diagnosis and cell typing, i
designed to detect single-nucleotide polymorphisms and cytosine
                                                                                                                                                                                                                                                                                                                                                                     Claim 1; SEQ ID NO 357813; 29pp + Sequence Listing; German.
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                                                                                                                              SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                            cancer; CNS;
primer; ss;
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                                                                                      Oligonucleotide primer SEQ ID NO 368590 for detecting SNP TSC0057099.
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Best Local Similarity 90.5
Matches 10; Conservative
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligoners for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligoners are also used for detecting cell type differentiation. ABC0010 represent the oligones described in the invention. ABC0010 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but
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Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                            Claim 1; SEQ ID NO 377538; 29pp + Sequence Listing; German.
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-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                         Sequence 12 BP; 4 A; 0 C; 3 G; S T; 0 U; 0 Other;
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99899, ABF00010-ABF99899, ABH00010-ABH99999 and ABI00010-ABB28073 represent the oligomers described in the invention. NoTE: The sequence data for this parent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                  SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                  Oligonucleotide primer SEQ ID NO 274417 for detecting SNP TSC0003540.
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              SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                              This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99999, ABF00010-ABF99999, ABH00010-ABH99999 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                 bet or oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                                                                                                                                                SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                            Gaps
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                                  Score 9.4; DB 1; Length 12; Pred. No. 1.2e+03; 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                         Oligonucleotide primer SEQ ID NO 311567 for detecting SNP TSC0024557.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic discorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABE99989, ABF00010-ABE99989, ABF00010-ABE99989, ABF00010-ABE99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo.int/pub/published_pot_sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                      Set of oligonucleotides, useful for diagnosis and cell typing, : designed to detect single-nucleotide polymorphisms and cytosine
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                                                                                                                                     Claim 1; SEQ ID NO 289935; 29pp + Sequence Listing; German.
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range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers ealso used for detecting cell type differentiation. ABC00010-ABC99989, ABC001010-ABF9989, ABH00010-ABH99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomers for detecting single nucleotides polymorphisms (SNP) oligomers are used for disquess and/or prognosis of cancer and range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC099889, ABC00010-ABE9989, ABH00010-ABH99989 and ABI00010-ABI82073 are represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at
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designed to detect single-nuclectide polymorphisms and cytosine
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designed to detect single-nucleotide polymorphisms and cytosine
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                                                                                                                                                             Berlin K;
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                                                                 07-APR-2000; 2000DE-01019173.
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Best Local Similarity 90.9'
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                                                                                                                                                                                                                                                                                designed to detect methylation status.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This invention describes novel oligonucleotide primers or peptide nucleic
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                                                                                       SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                             Oligonucleotide primer SEQ ID NO 348676 for detecting SNP TSC0000619.
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                                               This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABC0010-ABF9989, ABH0010-ABF9989 and ABI0010-ABI22073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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designed to detect single-nucleotide polymorphisms and cytosine
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                            Claim 1; SEQ ID NO 374901; 29pp + Sequence Listing; German.
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data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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WO200177384-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                            This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nuclectide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                    SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS;
                                                                                                               Oligonucleotide primer SEQ ID NO 267809 for detecting SNP TSC0000567.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Oligonucleotide primer SEQ ID NO 293415 for detecting SNP TSC0015609.
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                                          ABH67832 standard; DNA; 12 BP.
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                    RESULT 1678
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peptide nucleic acid, cytosine methylation, cardiovascular, primer, ss, central nervous system, gastrointestinal, respiratory, immune, metabolic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       was obtained in electronic format from Wl
ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                                                                                           This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, aradiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF9989, ABH00010-ABH99999 and ABI00010-ABI82073 tepresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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  (EPIG-) EPIGENOMICS AG
                                                                                                               WPI; 2001-657177/75
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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TAATTTTCTTT 12
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ABH7328/C
ID ABH7323/C
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AC ABH7232
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WO200177384-A2.
                                           18-OCT-2001.
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Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status. WPI; 2001-657177/75.

Berlin

Piepenbrock C,

olek A,

(EPIG-) EPIGENOMICS AG

German. Claim 1; SEQ ID NO 272307; 29pp + Sequence Listing; This invention describes novel oligonuclectide primers or peptide nucleic

Sequence 12 BP; 2 A; 0 C; 2 G; 8 T; 0 U; 0 Other;

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acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory. Central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99889, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI32073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at the printed specification, but ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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ABI06717:
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                                                                                                                                                                                                                                   SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                              Oligonucleotide primer SEQ ID NO 306319 for detecting SNP TSC0021944.
                                 Gaps
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          Score 9.4; DB 1; Length 12; Pred. No. 1.2e+03; 0; Mismatches 1; Indels
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                                                                                                                                       ABI06346 standard; DNA; 12 BP.
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                                                                                                                   SNP, single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                           Oligonucleotide primer SEQ ID NO 306690 for detecting SNP TSC0022131
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                                                                                                                                                                                                                                                                                                                                                                                                                                     06-APR-2001; 2001WO-IB000713.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PMA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99999, ABF00010-ABF9989, ABH00010-ABH99999 and ABI00010-ABI82073 tepresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                 oligonucleotides, useful for diagnosis and cell typing, ied to detect single-nucleotide polymorphisms and cytosine
                                                                                                                                                                                                                                                                                                                                              German.
                                                                                                                                                                                                                                                                                                                                           Claim 1; SEQ ID NO 306866; 29pp + Sequence Listing;
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                                                                          06-APR-2001; 2001WO-IB000713
                                                                                                               07-APR-2000; 2000DE-01019173
                                                                                                                                                                                        Piepenbrock C,
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WO200177384-A2
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Gaps ; 0 12.9%; Score 9.4; DB 1; Length 12; 90.9%; Pred. No. 1.2e+03; ive 0; Mismatches 1; Indels Local Similarity 90.9 946 GGTTTAATGTA 956 12 GGATTAATGTA 2 Matches g ò

AB136866/C

ID AB136866,
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AC AB136866;
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AC AB136866;
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DT 22-FEB-2002 (first entry)
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Oligonucleotide primer SEQ ID NO
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NOP; single nucleotide polymorph:
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NOPCHICAL INCOMES SYSTEM; gastroini
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Homo sapiens.
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NOC01777

Oligonucleotide primer SEQ ID NO 336839 for detecting SNP TSC0039548

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Berlin K;

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and oytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABE09989, ABF00010-ABE9989, ABF00010-ABE99989, ABF00010-ABE99989 and ABI00010-ABE82073 represent the oligomers described in the invention. NOTE: The sequence was obtained in electronic format from WIPD at the printed specification, but the wipo.int/pub/published_pct_sequences Gaps įs Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine methylation status. .. 0 Claim 1; SEQ ID NO 336839; 29pp + Sequence Listing; German. 12.9%; Score 9.4; DB 1; Length 12; 90.9%; Pred. No. 1.2e+03; ive 0; Mismatches 1; Indels Sequence 12 BP; 8 A; 2 C; 0 G; 2 T; 0 U; 0 Other; 10; Conservative 917 11 ATTTTTTGG 1 907 ATTTTCTTTGG Query Match Best Local Similarity Matches ઠ g

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ABI15225 standard; DNA; 12 BP. RESULT 1687

(first entry) 22-FEB-2002

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ABI15225;

Oligonucleotide primer SEQ ID NO 315198 for detecting SNP TSC0026773.

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Homo sapiens

WO200177384-A2.

18-OCT-2001.

06-APR-2001; 2001WO-IB000713

07-APR-2000; 2000DE-01019173

(EPIG-) EPIGENOMICS AG.

× Berlin Dlek A, Piepenbrock C,

WPI; 2001-657177/75.

Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.

claim 1; SEQ ID NO 315198; 29pp + Sequence Listing; German.

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The

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945 TGGTTTAATGT 955
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2 TGGATTAATGT 12
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oligomers are also used for detecting cell type differentiation. ABC00010 -ABC39989, ABF00010-ABF99889, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at flow. The bublished_pot_sequences
                                                                                                                                                                                                                                                                                                                                                                                           SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                                Oligonucleotide primer SEQ ID NO 291635 for detecting SNP TSC0014869.
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Matches 10; Conservative
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF9989, ABH00010-ABH9989 and ABI00010-ABI32073 tepseent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                                                                                                                                                                                              SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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ABI43225 standard; DNA; 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, contral nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF9989, ABH0010-ABH99989 and ABI00010-ABI82073 the represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                              SNP, single nucleotide polymorphism, human, diagnosis, PNA, cancer, CNS, peptide nucleic acid, cytosine methylation, cardiovascular, primer, ss; central nervous system; gastrointestinal, respiratory, immune, metabolic.
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Oligonucleotide primer SEQ ID NO 353162 for detecting SNP TSC0048346.
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Matches 10; Conservative
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABE99989, ABF00010-ABE99989, ABF00010-ABE99989, ABF00010-ABE99989 and ABI00010-ABE82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic formmat from WIPO at
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07-APR-2000; 2000DE-01019173
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les 10; Conservative
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                                             This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ACC0010 ABC99389, ABF00010-ABF99989, ABH00010-ABF99989 and ABI00010-ABF8073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Oligonucleotide primer SEQ ID NO 377568 for detecting SNP TSC0062396.
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Claim 1; SEQ ID NO 359703; 29pp + Sequence Listing; German.
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Best Local Similarity 90.9
Matches 10; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                   Gaps
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                                       Score 9.4; DB 1; Length 12; Pred, No. 1.2e+03;
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/*tag= a
/label= N-terminal of penicillinase
Sequence 12 BP; 8 A; 0 C; 2 G; 2 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                  5' end of penicillinase gene in plasmid pENX606.
                                                             Pred. No. 1.2e
0; Mismatches
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                                       12.9%;
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AAN50121 standard; DNA; 13 BP.
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84JP-00203772,
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                                                                                                                                                                                                                                                                                                                                                      (revised)
(first entry)
                                     Query Match
Best Local Similarity 90.9
Matches 10; Conservative
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Matches 10; Conservative
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P-PSDB; AAP50110.
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17-OCT-1991
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PCR primer; chromosomal abnormality; abnormality detection; leukaemia; lymphoma; carcinoma; adenocarcinoma; sarcoma; glioma; neuroblastoma; medullablastoma; malignant melanoma; malignant neoplastic condition; ss.

97WO-DK000556. 96DK-00001401.

08-DEC-1997;

06-DEC-1996;

Homo sapiens

Synthetic.

WO9824928-A2 11-JUN-1998. Pallisgaard N, Hokland

WPI; 1998-333344/29.

PALL/) PALLISGAARD N.

Primer ALL1:417L13 for abnormality detection.

25-SEP-1998 (first entry)

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Method for comparing mRNA from different nucleic acid samples - by reverse transcription and amplification using oligo-T primers.
                                                                                                                                                                          Parimoo S, Prouty SM, Stenn KS;
                                                               Enhanced specificity anchor primer, polyA tail; gene expression difference; cell type; ss.
                                                                                                                                                            (JOHJ ) JOHNSON & JOHNSON CONSUMER PROD
                                                  Enhanced specificity anchor primer 47.
                                                                                                                                                                                                                              Disclosure; Fig 4B; 44pp; English
       AAV03420 standard; DNA; 13 BP
                                                                                                                                 97WO-US005814.
                                                                                                                                               96US-0014666P.
                                    17-APR-1998 (first entry)
                                                                                                                                                                            Combates N, Pardinas JR,
                                                                                                                                                                                         WPI; 1997-503123/46.
                                                                                                     WO9737045-A1
                                                                                                                                 02-APR-1997;
                                                                                                                  09-OCT-1997.
                                                                                      Synthetic.
                      AAV03420;
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Detection of chromosomal abnormalities - by subjecting patient sample nucleic acids to a multiplex molecular amplification procedure using primers specific for characteristic nucleic acid sequence.

Claim 73; Page 67; 126pp; English.

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the polyA tail of mRNA and cDNA. The primers are of the general formula:

the polyA tail of mRNA and cDNA. The primers are of the general formula:

TIZNNIW, where M is A, G or C and N is A, G, C or T TABNIW, where we can be small to the compares the presence or in the method of the invention. This method compares the presence or level of individual mRNA molecules in at least 2 nucleic acid samples.

The method comprises contacting each of the nucleic acid samples with a colispodeoxynucleotide primer that hybridises to acid samples with a colispodeoxynucleotide primer that hybridises to a first site in mRNAs in the nucleic acid samples, reverse transcribing the mRNAs to which the primer hybridises to produce a population of DNA strands that are complementary to the mRNAs in the 2 samples. The amount of cDNA produced is quantified. The populations of cDNA are contacted with a second oligodeoxynucleotide primer (e.g. present primer) that hybridises to a second site in the cDNA populations, the contact being performed under conditions in which the second populations of the DNA strands are subcond population of the DNA strands are second populations of amplification products in the first and second populations of amplification products in the first and second con be used for screening differences in gene expression between cell types or between cells in different stages of development or various cell types or between cells in different stages of development or various cell types or between cells in different stages of development or various cell types or between cells in different stages of development or various cell types or between cells in different stages of development or various cell types or between cells in different stages of development or various cell types or between cells in different stages of development or various cell types or between cells in different stages of development or various cell types or between cells and conditions
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Pred. No. 1.3e+03;
0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    12.9%;
90.9%;
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Best Local Similarity 90.9
Matches 10; Conservative
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                                                                                                                                                                                                                                                 associated with diseases including numerous leukaemia's, lymphoma's, carcinoma's, adenocarcinoma's, sarcoma's, glioma's, neuroblastoma's, medullablastoma, malignant melanoma, and malignant neoplastic conditions
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0
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Pred. No. 1.3e+03;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                     Sequence 13 BP; 1 A; 2 C; 3 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                         0; Mismatches
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                                                                                                                                                                                                                                                                                          10; Conservative
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Best Local Similarity
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BP.

AAV40929 standard; DNA; 13

AAV40929 RESULT 1696 AAV40929 ID AAV4092 XX AC AAV4092X

908 TTTTCTTTGGT 918

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Stephens JC;

Nandabalan K,

Duda A,

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Denton RR,
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                                       Human Ostec
nucleotide
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 Chew A,
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Matches
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                                                                                                                                                                                                                                                  The present invention describes a method for the species-specific detection of a Trichosporon genus microbe which includes detecting a polymuclectide specific to the species of a Trichosporon genus microbe. Trichosporon polymuclectides can be used for the diagnosis and treatment of Trichosporosis. The method can distinguish Trichosporosis species to species level rapidly in high precision. AAAA6734 to AAA26849 represent polymuclectide sequences from various Trichosporon species, which are
                                                                                                                                                                                           ಹ
                                                                                                                                                                                       Species-specific detection of a Trichosporon genus microbe species and new polynucleotide - used for the diagnosis and the treatment of Trichosporosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; TNFRSF11E; osteoclastogenesis inhibitory factor; single nucleotide polymorphism; SNP; osteoclast recruitment; osteoclast function; osteocorosis; metastatic bone disease; Paget's disease; rheumatoid arthritis; periodontal bone disease; ASO; allele-specific oligonucleotide; probe; ss.
                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                   Trichosporon genus microbe; detection; species-specific; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                   ;
0
Trichosporon aquatile polynucleotide sequence SEQ ID NO:62
                                                                                                                                                                                                                                                                                                                                                            Score 9.4; DB 1; Length 13;
Pred. No. 1.3e+03;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                      used in the exemplification of the present invention
                                                                                                                                                                                                                                                                                                                                           Sequence 13 BP; 5 A; 2 C; 2 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human TNFRSF11B gene ASO probe, SEQ ID NO: 112.
                                                                                                                                                                                                                               Claim 2; Page 40; 47pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (GENA-) GENAISSANCE PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAF70056 standard; DNA; 13 BP
                                                                                                                                                                                                                                                                                                                                                              12.9%;
                                                                                                           98JP-00237060,
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                                                                                                                              98JP-00237060
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                                                                                                                                                                                                                                                                                                                                                             Query Match 12.9
Best Local Similarity 90.9
Matches 10; Conservative
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                                                                                                                                                (IATR ) IATRON LAB INC
                                                Trichosporon aquatile.
                                                                                                                                                                   WPI; 2000-249679/22.
                             trichosporosis; ds
                                                                  JP2000060564-A.
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                                                                                                           24-AUG-1998;
                                                                                                                              24-AUG-1998;
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                                                                                       29-FEB-2000
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AAF70056
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                                                                       Osteoclastogenesis Inhibitory Factor nucleotides, comprising single otide polymorphisms, useful for studying e.g. osteoporosis, Paget's
                                                                                                                                                                                                                                                                                                the human
                                                                                                                                                                                                                                                                                    The present sequence is a probe used to detect polymorphisms in the human osteoclastogenesis inhibitory factor (TNFRSF11B). Polymucleotides comprising one or more of twenty four novel single nucleotide polymorphisms in the TNFRSF11B gene have been identified. TNFRSF11B regulate osteoclast recruitment and function. An understanding of variations in the gene should thus be useful in developing new therapies for metabolic disorders caused by abnormal osteoclast recruitment and function such as osteoporosis, metastatic bone disease, Paget's disease, rheumatoid arthritis and periodontal bone disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ·;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12.9%; Score 9.4; DB 1; Length 13; ilarity 90.9%; Pred. No. 1.38+03; Conservative 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 13 BP; 1 A; 4 C; 2 G; 6 T; 0 U; 0 Other;
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                                                                                                                                                                                                                     Claim 15; Page 23; 114pp; English
                                                                                                                                                 disease and rheumatoid arthritis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP.
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WPI; 2001-147175/15
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905 TCATTITCTT 915

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oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99889, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NoTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, contral nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Oligonucleotide SEQ ID NO 21609 for detecting SNP TSC0004336.
                                                                                                                                                                           12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03; tive 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; SEQ ID NO 21609; 29pp + Sequence Listing; German.
                                                                                                                                          Sequence 13 BP; 4 A; 3 C; 1 G; 4 T; 0 U; 1 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 13 BP; 7 A; 0 C; 3 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                     ABC21592 standard; DNA; 13
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                                                                                                                                                                                                                         10; Conservative
                                                                                                                                                                                                                                                               955 TATCGCTACCA 965
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                                                                                                                                                                                                                                                                                                   3 TATCGCTATCA 13
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Best Local Similarity
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                                                                                                                                                                                                       SNP, single nucleotide polymorphism; human; diagnosis; PNA, cancer; CNS, peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                                                            Oligonucleotide SEQ ID NO 23962 for detecting SNP TSC0005553.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; SEQ ID NO 23962; 29pp + Sequence Listing; German.
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                                                                               ABC23945 standard; DNA; 13
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13 ATTTGTTTTGGTY 1
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13 rcacrrrcrr
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                                                RESULT 1701
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21-FEB-2002 (first entry)

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ABC49345;

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and merabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABE99989, ABF00010-ABE99989, ABF00010-ABE99989, ABF00010-ABE99989, ABF00010-ABE99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at the printed specification, but fire wipo int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SNP, single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Oligonucleotide SEQ ID NO 51424 for detecting SNP TSC0014354.
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                                                                                                                                                                                                                                                                                            Claim 1; SEQ ID NO 51054; 29pp + Sequence Listing; German.
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                   07-APR-2000; 2000DE-01019173
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Best Local Similarity 90.5
Matches 10, Conservative
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TTGGTTTATTG 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99999, ABF00010-ABF99999, ABH00010-ABH99999 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                           SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Oligonucleotide SEQ ID NO 49362 for detecting SNP TSC0013972
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12.9%; Score 9.4; DB 1; Length 13;
Best Local Similarity 90.9%; Pred. No. 1.3e+03;
Matches 10; Conservative 0; Mismatches 1; Indels
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Sequence 13 BP; 8 A; 1 C; 3 G; 1 T; 0 U; 0 Other;

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99999, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI32073 tepresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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designed to detect single-nucleotide polymorphisms and cytosine
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Claim 1; SEQ ID NO 51424; 29pp + Sequence Listing; German.
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acid (PNA) oligomers for detecting single nucledide polymorphisms (SNP) and cytosine methylation status in chemically pretrated genomic DNA. The oligomucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at

This invention describes novel oligonucleotide primers or peptide nucleic

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                                                                                                                                                                                                                                                                                                                    SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Score 9.4; DB 1; Length 13;
Pred. No. 1.3e+03;
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RESULT 1707 ABC54442

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and merabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at fire wipo int/pub/published_pct_sequences
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                                 Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                          SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                               Oligonucleotide SEQ ID NO 54459 for detecting SNP TSC0014930.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 1; SEQ ID NO 54459; 29pp + Sequence Listing; German.
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Matches 10, Conservative
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Pred. No. 1.3e+03;
1; Mismatches 2; Indels
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                                                                                                                               onucleotides, useful for diagnosis and cell detect single-nucleotide polymorphisms and
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Best Local Similarity 76.9%;
Matches 10; Conservative
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                                                                                                                                                              designed to detect methylation status.
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oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at figurial form.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Sequence 13 BP; 5 A; 0 C; 6 G; 2 T; 0 U; 0 Other;

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acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 the septemble for this patent did not form part of the printed specification, but was obtained in electronic formet from WIPO at
                                                                                                    SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                             Oligonucleotide SEQ ID NO 16207 for detecting SNP TSC0003545.
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                                                                                                                                                                                                                                                                                                                                                                                 SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Pred. No. 1.3e+03;
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ftp.wipo.int/pub/published_pct_sequences
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Matches 10; Conservative
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Set of oligonucleotides, useful for diagnosis and cell typing, is

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Piepenbrock C,

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                                                                                                                                                             This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99999, ABF00010-ABF99999, ABH00010-ABH99999 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                          Claim 1; SEQ ID NO 119572; 29pp + Sequence Listing; German
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Matches 10; Conservative
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, contral nervous system, ardiovascular and metabolic disorders. The coligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                     SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                                          Score 9.4; DB 1; Length 13; Pred. No. 1.3e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SNP; single nuclectide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Pred. No. 1.3e+03;
0; Mismatches 1; Indels
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99889, ABF00010-ABF99889, ABH00010-ABH99989 and ABI00010-ABF3073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                                                                                                 Claim 1; SEQ ID NO 173549; 29pp + Sequence Listing; German.
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretraeted genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99899, ABF00010-ABF99899 ABH00010-ABH99989 and ABI00010-ABI82073 tepresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo.int/pub/published_pct_sequences
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RESULT 1724 ABF75595/c WO200177384-A2.

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPD at
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, contral nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABT00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from MIPO at
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disonders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99899, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but typo.int/pub/published_pct_sequences
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ABH03395 standard; DNA; 13
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Oligonucleotide SEQ ID NO 205979 for detecting SNP TSC0050473.

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                                                                                                                                                                                                                                             SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                              Oligonucleotide SEQ ID NO 229628 for detecting SNP TSC0056011.
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943 ATTGGTTTAAT 953
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tes 10; Conserv
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22-FEB-2002 (first entry)

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                                         SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00110 absorpes, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                     Claim 1; SEQ ID NO 232882; 29pp + Sequence Listing; German.
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                                                                                                                                                                                                                                                                                                                       This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99999, ABF00010-ABE99999, ABH00010-ABH99999 and ABI00010-ABH32073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but ftp.wipo.int/pub/published_pct_sequences
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06-APR-2001; 2001WO-IB000713
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Oligonucleotide SEQ ID NO 184801 for detecting SNP TSC0045589.
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                                                                                                                     Query Match 12.9%; Score 9.4; DB 1; Length 13; Best Local Similarity 90.9%; Pred. No. 1.3e+03; Matches 10; Conservative 0; Mismatches 1; Indels
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                                                                       Sequence 13 BP; 9 A; 3 C; 0 G; 1 T; 0 U; 0 Other;
was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                     SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Piepenbrock C,

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, contral nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at
central nervous system; gastrointestinal; respiratory; immune; metabolic.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC09989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                             ligonucleotides, useful for diagnosis and cell typing, : to detect single-nucleotide polymorphisms and cytosine
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                                                                                                                                                  Claim 1; SEQ ID NO 163722; 29pp + Sequence Listing; German
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and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF0010-ABF99999, ABF00010-ABF99999 and ABI00010-ABI82073 represent the oligomers described in the invention. NoTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Pred. No. 1.3e+03;
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                                                                                                                                                                                                                                                                                                     SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Pred. No. 1.3e+03;
0; Mismatches 1; Indels
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Score 9.4; DB 1;
Pred. No. 1.3e+03;
                              1; Mismatches
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                                                                946 GGTTTAATGTATC 958
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ABH58823 standard; DNA; 13 BP.

ABH58823/c ID ABH588 XX

Sequence 13 BP; 2 A; 0 C; 2 G; 8 T; 0 U; 1 Other;

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This invention describes novel oligonucleotide primers or peptide nucleic acid (FNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                                                              Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine methylation status.
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Best Local Similarity 90.9
Matches 10; Conservative
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ABH59591/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but fur wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                 SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                       Oligonucleotide SEQ ID NO 258800 for detecting SNP TSC0062902.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but two was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SNP; single nuclectide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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onucleotides, useful for diagnosis and cell typing, idetect single-nucleotide polymorphisms and cytosine
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                                                                 Claim 1; SEQ ID NO 259568; 29pp + Sequence Listing; German.
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                 designed to detect methylation status.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF9989, ABH00010-ABH99989 and ABT00010-ABI82073 trepresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from NIPO at
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                represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073
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.larity 76.9%; Pred. No. 1.3e+03;
Conservative 1; Mismatches 2; Indels
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12.9%; Score 9.4; DB 1; Length 13;
Best Local Similarity 76.9%; Pred. No. 1.38+03;
Matches 10; Conservative 1; Mismatches 2; Indels
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                                                                                                                 Sequence 13 BP; 5 A; 4 C; 0 G; 3 T; 0 U; 1 Other;
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946 GGTTTAATGTATC 958

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Matches

This invention describes novel oligonucleotide primers or peptide nucleic acid (PMA) oligomers for detecting single nucleotide polymorphisms (SMP) and cytosine methylation status in chemically pretreated genomic DMA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABCO0010

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WPI; 2001-657177/75.
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                                                                                                                                                    SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                Oligonucleotide SEQ ID NO 93489 for detecting SNP TSC0023360.
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                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; SEQ ID NO 93489; 29pp + Sequence Listing; German.
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Best Local Similarity 76.9%;
Matches 10; Conservative
                                                                ABC93472 standard; DNA; 13
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(EPIG-) EPIGENOMICS AG
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                                                                                      ABC93472;
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genemic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 1; SEQ ID NO 94714; 29pp + Sequence Listing; German.
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Best Local Similarity 90.9
Matches 10; Conservative
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schultz1-899.rng

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF99989, ABH00010-ABF99989 and ABI00010-ABF8073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at

Berlin K;

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Gaps

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Query Match

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946 GGTTTAATGTA 956

12 GGTTTATTGTA 2

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Sequence 13 BP; 6 A; 4 C; 0 G; 3 T; 0 U; 0 Other;

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligoners for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligoners are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at the printed specification, but fire wipo.int/pub/published_pct_sequences
                                                                                                        oligonuclectides, useful for diagnosis and cell typing, is to detect single-nuclectide polymorphisms and cytosine
                                                                                                                                                           Claim 1; SEQ ID NO 71612; 29pp + Sequence Listing; German.
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07-APR-2000; 2000DE-01019173.
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                                                    Piepenbrock C,
                          (EPIG-) EPIGENOMICS AG
                                                                              WPI; 2001-657177/75
                                                                                                                       designed to detect methylation status.
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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BP.
ABC21785 standard; DNA; 13
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Oligonucleotide SEQ ID NO 98934 for detecting SNP TSC0024573.

(first entry)

21-FEB-2002

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2, Indels

12.9%; Score 9.4; DB 1; Length 13; 76.9%; Pred. No. 1.3e+03;

1; Mismatches

ABC98917;

BP

ABC98917 standard; DNA; 13

RESULT 1750

ABC989

Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine Claim 1; SEQ ID NO 98934; 29pp + Sequence Listing; German. methylation status.

Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.

Berlin K;

Olek A, Piepenbrock C,

WPI; 2001-657177/75

(EPIG-) EPIGENOMICS AG.

06-APR-2001; 2001WO-IB000713 07-APR-2000; 2000DE-01019173 Claim 1; SEQ ID NO 21802; 29pp + Sequence Listing; German

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Piepenbrock C,

olek A,

(EPIG-) EPIGENOMICS AG

06-APR-2001; 2001WO-IB000713 07-APR-2000; 2000DE-01019173

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RESULT 1751

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acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABR59989, ABR0010-ABR99989 and ABI0010-ABR82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at the printed specification, but fire, wipo.int/pub/published_pot_sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This invention describes novel oligonucleotide primers or peptide nucleic
                                                                                                                                                                                              SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                       ABC50399 standard; DNA; 13
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RESULT 1753
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                               Score 9.4; DB 1; Length 13
Pred. No. 1.3e+03;
1; Mismatches 2; Indels
                         Sequence 13 BP; 6 A; 4 C; 0 G; 2 T; 0 U; 1 Other;
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Matches 10; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                       Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                           Claim 1; SEQ ID NO 28627; 29pp + Sequence Listing; German.
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ABC31427 standard; DNA; 13
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les 10; Conservative
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range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABF00010-ABF99989, ABF00010-ABF99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at fire, wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Matches 10, Conservative
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Length 13;

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Query Match Best Local Similarity

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                                                                                                                                                                                                                                                                                                     peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Mismatches
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF99989, ABH0010-ABF99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                                                                                                 Claim 1; SEQ ID NO 109660; 29pp + Sequence Listing; German.
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                                                                                   SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; 8s; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                        Oligonucleotide SEQ ID NO 109123 for detecting SNP TSC0027313.
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                                                                                                             This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF99899, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo.int/pub/published_pct_sequences
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                                                           Claim 1; SEQ ID NO 61983; 29pp + Sequence Listing; German
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                                                                                                                                                                                                                                                                                                                                                                                                                                     SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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designed to detect single-nucleotide polymorphisms and cytosine
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                                                                                                                           SNP; single nuclectide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                         Oligonucleotide SEQ ID NO 39917 for detecting SNP TSC0012171
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acid; cytosine methylation; cardiovascular; primer; 88; system; gastrointestinal; respiratory; immune; metabolic.
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                                                                  set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99889, ABF00010-ABF99989, ABH00010-ABF99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at
acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically precreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory. Central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABC00010-ABC9989, ABC00010-ABC99989, ABC00010-ABC99989, ABC00010-ABC99989, ABC00010-ABC99989, ABC00010-ABC99989, ABC00
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                      Oligonucleotide SEQ ID NO 137356 for detecting SNP TSC0034314.
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                                                                                                                                                                                                                                                                                                    SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                             Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                                                                                                                                                        Oligonucleotide SEQ ID NO 127229 for detecting SNP TSC0031843.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; SEQ ID NO 127229; 29pp + Sequence Listing; German.
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           Score 9.4; DB 1; Length 13; Pred. No. 1.3e+03; 0; Mismatches 1; Indels
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90.9%;
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                             Best Local Similarity 90.9
Matches 10; Conservative
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nes 10; Conservative
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                Query Match
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Α, Berlin

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                                              This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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Claim 1; SEQ ID NO 137356; 29pp + Sequence Listing; German.
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Homo sapiens.

ABF37359 standard; DNA; 13

RESULT 1769

ABF37359/ ID ABF3

942 CATTGGTTTAA 952

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Matches

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ABH21913;
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                                                                                                                                                                                                            This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99999, ABF00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at the printed specification, but fire wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                       Berlin K;
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Best Local Similarity 90.9
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                                                                                                       Olek A, Piepenbrock C,
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TCACTACCAAC 1
                                                                                  (EPIG-) EPIGENOMICS AG
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WO200177384-A2
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic discorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABE09989, ABF00010-ABE9989, ABH0010-ABE9989, ABH0010-ABE9989, and ABI00010-ABE32073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                                                               Claim 1; SEQ ID NO 219390; 29pp + Sequence Listing; German.
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oligomers are also used for detecting cell type differentiation. ABC00010 -ABC99889, ABF00010-ABF99889, ABF00010-ABF99889, ABF00010-ABF99889, ABF00010-ABF99889, ABF00010-ABF89989 and ABI00010-ABF88073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                        SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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956 ATCGCTACCAA 966
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                               3 Arctcraccaa 13
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF9989, ABH0010-ABH99999 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but
                                 SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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designed to detect single-nuclectide polymorphisms and cytosine
Oligonuclectide SEQ ID NO 226464 for detecting SNP TSC0055199,
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABE99999, ABF00010-ABE99999, ABF00010-ABE99999, abmoorle in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, aradiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                   This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastroincestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The coligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99899, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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Claim 1; SEQ ID NO 154247; 29pp + Sequence Listing; German.
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                                                                                                                                                                                                                                                                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                               Score 9.4; DB 1; Length 13; Pred. No. 1.3e+03; 0; Mismatches 1; Indels
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Sequence 13 BP; 4 A; 5 C; 0 G; 4 T; 0 U; 0 Other;
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                                                                                                                 SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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designed to detect single-nucleotide polymorphisms and cytosine
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                                                                                           Oligonucleotide SEQ ID NO 208535 for detecting SNP TSC0050953.
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC09989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the publypublished_pot_sequences
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WO200177384-A2
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                                                             Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                       Claim 1; SEQ ID NO 184805; 29pp + Sequence Listing; German.
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Pred. No. 1.3e+03;
0; Mismatches 1; Indels
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF9989, ABH00010-ABH9989 and ABT00010-ABI82073 tepses the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from MIPO at
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12.9%; Score 9.4; DB 1; Length 13;

Query Match

RESULT 1785 ABH12346

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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                               SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                       Oligonucleotide SEQ ID NO 241734 for detecting SNP TSC0058949.
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12.9%; Score 9.4; DB 1; Length 13;
Best Local Similarity 76.9%; Pred. No. 1.3e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; SS; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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to detect single-nucleotide polymorphisms and cytosine
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                                                                                                                                                                                 Claim 1; SEQ ID NO 242136; 29pp + Sequence Listing; German.
                                                                                                                                                                                                                                                                                                                                                                       Score 9.4; DB 1; Length 13;
Pred. No. 1.3e+03;
0; Mismatches 1; Indels
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                                                                                         Berlin K;
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                       06-APR-2001; 2001WO-IB000713.
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Matches 10; Conservative
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                                                                                                                                                           methylation status.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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Claim 1; SEQ ID NO 245087; 29pp + Sequence Listing;
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABR00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from MIPO at
                                                                                                                                                                                                                                             SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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represent the oligomers described in the invention. NOTE: The sequence data for this parent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                             Query Match
12.9%; Score 9.4; DB 1; Length 13;
Best Local Similarity 76.9%; Pred. No. 1.3e+03;
Matches 10; Conservative 1; Mismatches 2; Indels
                                                                                           Sequence 13 BP; 3 A; 0 C; 3 G; 6 T; 0 U; 1 Other;
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Matches 10; Conservative
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                                                                            12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03; ive 0; Mismatches 1; Indels
                                             Sequence 13 BP; 2 A; 0 C; 3 G; 8 T; 0 U; 0 Other;
was obtained in electronic format from Wittp.wipo.int/pub/published_pct_sequences
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                         Claim 1; SEQ ID NO 261561; 29pp + Sequence Listing; German.
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Best Local Similarity
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                                                Homo sapiens.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABF99989, ABH00010-ABF99989, and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                                                                                                                                                                                                                                                                               Claim 1; SEQ ID NO 45664; 29pp + Sequence Listing; German.
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                                                                      Berlin K;
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Sequence 13 BP; 9 A; 3 C; 0 G; 1 T; 0 U; 0 Other;

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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nuclectide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99899, ABF00010-ABF99899, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at the printed specification, but fire wipo.int/pub/published_pct_sequences
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acid (PNA) oligomers for detecting single nuclectide polymorphisms (SNP) and oytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting call type differentiation. ABC0010-ABC09989, ABC0010-ABE09989, ABF00010-ABE99989, ABF00010-ABE99989 and ABI00010-ABE82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP)
                                                                                                                                  SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory. Central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99889, ABF00010-ABE99889, ABF00010-ABE99889 and ABI00010-ABE3073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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Best Local Similarity 90.9
Matches 10; Conservative
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                                                   methylation status.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretraeted genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, contral nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99899, ABF00010-ABF99899, ABH00010-ABH99989 and ABT00010-ABI82073 trepresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from MIPO at
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central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00110 -ABC9989, ABF00010-ABF9989, ABF00010-ABF9989 and ABI00010-ABF8073 represent the oligomers described in the invention. NoTE: The sequence adar for this parent did not form part of the printed specification, but was obtained in electronic format from WIPO at the printed specification, but ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                                                                                                     12.9%; Score 9.4; DB 1; Length 13; 76.9%; Pred. No. 1.3e+03; ive 1; Mismatches 2; Indels
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Best Local Similarity 76.9'
Matches 10, Conservative
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Oligonucleotide SEQ ID NO 31443 for detecting SNP TSC0009724.

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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nuclectide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABE99989, ABF00010-ABE99989 and ABI00010-ABE3073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from wIPO at
                                                                                                                                                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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903 GGTCATTTTCTTT
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo int/pub/published_pct_sequences
                                             SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Matches 10; Conservative
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ò 셤 Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.

WPI; 2001-657177/75.

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic formmat from WIPO at
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 9.4; DB 1; Length 13;
Pred. No. 1.3e+03;
0; Mismatches 1; Indels
Claim 1; SEQ ID NO 88331; 29pp + Sequence Listing; German.
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90.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. AGC00010-ABC99989, ABF00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                           SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                          Gaps
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                                                                                                                                                                                                                                                                                    Oligonucleotide SEQ ID NO 63680 for detecting SNP TSC0016816.
                                                            12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1; SEQ ID NO 63680; 29pp + Sequence Listing; German.
                                      Sequence 13 BP; 1 A; 1 C; 4 G; 7 T; 0 U; 0 Other;
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was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                          ABC63663 standard; DNA; 13 BP.
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                                                                           Best Local Similarity 90.9
Matches 10, Conservative
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                                                              Query Match
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                              SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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ftp.wipo.int/pub/published_pct_sequences
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ABC39010 standard; DNA; 13
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                                                                                                                    20-FEB-2002
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Gaps

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12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03; ive 0; Mismatches 1; Indels

Query Match Best Local Similarity 90.9 Matches 10; Conservative

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF99989, ABH00010-ABF99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic formmat from WIPO at
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cycosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                       Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                                                                                     Claim 1; SEQ ID NO 64546; 29pp + Sequence Listing; German
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                   Berlin K;
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                      Piepenbrock C,
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central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Best Local Similarity 90.5-
Fines 10; Conservative
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                                            Homo sapiens.
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RESULT 1810 ABC64529

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Gaps

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Gaps

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Score 9.4; DB 1; Length 13; Pred. No. 1.3e+03; 0; Mismatches 1; Indels

12.9%; 90.9%;

Query Match
Best Local Similarity 90.>
Local Similarity 90.>
Local Similarity 90.>

907 ATTTTCTTTGG 917

12 ATTTTTTTGG

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and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF9989, ABF00010-ABF9989, ABF00010-ABF9989, and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but they was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99889, ABH00010-ABH99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPD at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                              1; Indels
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90.9%;
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Best Local Similarity 90.9
Matches 10; Conservative
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Berlin K;

Piepenbrock C,

olek A,

(EPIG-) EPIGENOMICS AG.

06-APR-2001; 2001WO-IB000713. 07-APR-2000; 2000DE-01019173

WO200177384-A2. Homo sapiens.

18-OCT-2001.

Oligonucleotide SEQ ID NO 115747 for detecting SNP TSC0029020.

BP.

ABF15750 standard; DNA; 13

RESULT 1813

ABF157

21-FEB-2002 (first entry)

ABF15750;

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Sequence 13 BP; 9 A; 2 C; 0 G; 2 T; 0 U; 0 Other;

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABE9989, ABF00010-ABF99989, ABH00010-ABF99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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ABF28323 standard; DNA; 13
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Best Local Similarity 90.9
Matches 10; Conservative
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                                                                                                                          SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                    Oligonucleotide SEQ ID NO 122101 for detecting SNP TSC0030522.
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Best Local Similarity 90.2
Best Local 10; Conservative
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ABF22104;
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Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                      Claim 1; SEQ ID NO 128320; 29pp + Sequence Listing; German.
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Pred. No. 1.3e+03;
1; Mismatches 2; Indels
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Best Local Similarity 76.9%;
Matches 10; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at the printed specification, ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                             Gaps
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                                                                                                                                         Score 9.4; DB 1; Length 13; Pred. No. 1.3e+03; 0; Mismatches 1; Indels
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PMA) oligoners for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010

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acid (FNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory. Central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99899, ABC0010-ABC99899 ABC0010-ABC99899 ABC0010-ABC9989 and ABI0010-ABCS073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo int/pub/published_pct_sequences
               SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                          onucleotides, useful for diagnosis and cell typing, 3 detect single-nucleotide polymorphisms and cytosine
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  designed to detect methylation status.
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Best Local Similarity
Matches 10; Conserv
                                                                                                                                          WO200177384-A2
                                                                                                      Homo sapiens
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                                                                                                                                                                                                                                                                                      SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                              Oligonucleotide SEQ ID NO 221889 for detecting SNP TSC0053997
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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designed to detect single-nucleotide polymorphisms and cytosine
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                                                                                     methylation status.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF0010-ABF9989, ABH0010-ABF9989, and ABI0010-ABI22073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic form at from MIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 13 BP; 7 A; 4 C; 0 G; 2 T; 0 U; 0 Other;
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ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC09989, ABF00010-ABF9989, ABH00010-ABH99899 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at the printed specification, but ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                         Berlin K;
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                                                                                                                                         Piepenbrock C,
                                                                        (EPIG-) EPIGENOMICS AG
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Gaps

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12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03; tive 0; Mismatches 1; Indels

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretraeted genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic discorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF9989, ABF00010-ABF99989 and ABI00010-ABF9989, represent the oligomers described in the invention. NOTE: The sequence and not this patent did not form part of the printed specification, but two obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                  SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                   Oligonucleotide SEQ ID NO 180827 for detecting SNP TSC0044744.
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                                                                                                                            ABF80830 standard; DNA; 13
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941 TCATTGGTTTA 951
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                     SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                               Oligonucleotide SEQ ID NO 208264 for detecting SNP TSC050910.
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BP.
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Conservative
ABH08287 standard; DNA; 13
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nes 10; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Pred. No. 1.3e+03;
0; Mismatches 1; Indels
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Sequence 13 BP; 5 A; 0 C; 2 G; 6 T; 0 U; 0 Other;
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                                     Query Match
Best Local Similarity 90.9%;
Matches 10; Conservative
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Matches 10; Conserv
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99889, ABF00010-ABF99899, ABF00010-ABF99989, ABF00010-ABF99989 and ABI00010-ABF8073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but typ.wipo.int/pub/published_pot_sequences
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                                              Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine
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designed to detect single-nucleotide polymorphisms and cytosine
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                                                                                                                 Claim 1; SEQ ID NO 237191; 29pp + Sequence Listing; German.
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            WPI; 2001-657177/75.
                                                                  designed to detect amethylation status.
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les 10; Conserv
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   Homo sapiens.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, aradiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                                                                                                                                                          12.9%; Score 9.4; DB 1; Length 13; 76.9%; Pred. No. 1.3e+03; tive 1; Mismatches 2; Indels
                                                                                                                                                                                                              Sequence 13 BP; 5 A; 1 C; 1 G; 5 T; 0 U; 1 Other;
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Length 13;

12.9%; Score 9.4; DB 1; 90.9%; Pred. No. 1.3e+03;

Query Match Best Local Similarity

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Pred. No. 1.3e+03;
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Mismatches
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ftp.wipo.int/pub/published_pct_sequences
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90.9%;
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Best Local Similarity 90.9
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18-OCT-2001

olek A,

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99999, ABF00010-ABF99999, ABH00010-ABH99999 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic formmat from WIPO at
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designed to detect single-nucleotide polymorphisms and cytosine
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                                                                                                                                                                                                                                                                                                                                                     claim 1; SEQ ID NO 243911; 29pp + Sequence Listing; German.
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Matches 10; Conserv
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                                                                                   SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                             Oligonucleotide SEQ ID NO 217201 for detecting SNP TSC0052794.
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Query Match

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Matches
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                                               This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, actdiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                          Claim 1; SEQ ID NO 245751; 29pp + Sequence Listing; German.
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data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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Local Similarity 90.9%; Pred. No. 1.3e+03; les 10; Conservative 0; Mismatches 1:
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                                                                                                                           SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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designed to detect single-nucleotide polymorphisms and cytosine
                                                                                                      Oligonucleotide SEQ ID NO 260385 for detecting SNP TSC0004827.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI32073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic formmat from WIPO at
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ö This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and oytosine methylation status in chemically pretreated genomic DNA. The oligomucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99889, ABF00010-ABF99889, ABH00010-ABH99889 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at This invention describes novel oligonucleotide primers or peptide nucleic SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic. 0; Gaps Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine methylation status. Oligonuclectide SEQ ID NO 92856 for detecting SNP TSC0023219. / Match 12.9%; Score 9.4; DB 1; Length 13; Local Similarity 90.9%; Pred. No. 1.3e+03; nes 10; Conservative 0; Mismatches 1; Indels Claim 1; SEQ ID NO 92856; 29pp + Sequence Listing; German Claim 1; SEQ ID NO 42401; 29pp + Sequence Listing; German. Sequence 13 BP; 7 A; 0 C; 3 G; 3 T; 0 U; 0 Other; Berlin K; Berlin 踞. 06-APR-2001; 2001WO-IB000713. 07-APR-2000; 2000DE-01019173 ABC92839 standard; DNA; 13 (first entry) 935 TCCTCTTCATT 945 Olek A, Piepenbrock C, Olek A, Piepenbrock C, (EPIG-) EPIGENOMICS AG 13 racrerrearr 3 EPIGENOMICS AG WPI; 2001-657177/75 WPI; 2001-657177/75 methylation status WO200177384-A2. Homo sapiens 18-OCT-2001. 21-FEB-2002 ABC92839; Query Match (EPIG-) RESULT 1839 Matches ABC92839, à 셤

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF99989, ABH00010-ABF9989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic formmat from WIPO at fitted specification, but fit.wipo.int/pub/published_pct_sequences
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acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and range of diseases including immune system, gastrointestinal, respiratory. central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99889, ABC0010-ABC99889 ABC0010-ABC99889 and ABL0010-ABC0010 ABC0010 ABC0
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to detect single-nucleotide polymorphisms and cytosine
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                                                                                                                                                                                                                                                                                                                                                                      Sequence 13 BP; 7 A; 4 C; 0 G; 2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Mismatches
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90.9%;
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Best Local Similarity 90.9
Matches 10; Conservative
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ABC95529/
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Oligonucleotide SEQ ID NO 21801 for detecting SNP TSC0004359.

20-FEB-2002 (first entry)

ABC21784;

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                                                                                                                                                                                                                                                                                                                             SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                  Gaps
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designed to detect single-nucleotide polymorphisms and cytosine
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               Length 13;
Score 9.4; DB 1; Lengtn 13, Pred. No. 1.3e+03;
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                                             1; Mismatches
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               12.9%;
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                                                                            920 TTTGCCTTTTATC 932
                                                                                                                                                                                                                                                              (first entry)
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13 TTTGTTTTTATY 1
                                               10; Conservative
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             Query Match
Best Local Similarity
Matches 10; Conserv
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Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine

X, Berlin

Olek A, Piepenbrock C,

WPI; 2001-657177/75.

methylation status.

(EPIG-) EPIGENOMICS AG

06-APR-2001; 2001WO-IB000713. 07-APR-2000; 2000DE-01019173

WO200177384-A2. Homo sapiens

18-OCT-2001.

Claim 1; SEQ ID NO 21801; 29pp + Sequence Listing; German.

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC009989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 are present the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            was obtained in electronic format from WI ftp.wipo.int/pub/published_pct_sequences
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Best Local Similarity 90.9
Matches 10, Conservative
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Homo sapiens

RESULT 1842 ABC21784 ID ABC21784 standard; DNA; 13 BP.

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Gaps

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Score 9.4; DB 1; Length 13; Pred. No. 1.3e+03;); Mismatches 1; Indels

Sequence 13 BP; 3 A; 3 C; 0 G; 7 T; 0 U; 0 Other;

12.9%; 90.9%;

Query Match Best Local Similarity 90.9 Matches 10; Conservative

TCATTTTCTTT 915

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                                                                                                                                                                                                                                                                                                                 This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretracted genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99889, ABF00010-ABF99889, ABH00010-ABH99889 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at the printed specification, but fry.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                         Claim 1; SEQ ID NO 97201; 29pp + Sequence Listing;
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                                                              06-APR-2001; 2001WO-IB000713.
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Best Local Similarity 90.9%;
Matches 10; Conservative
                                                                                             07-APR-2000; 2000DE-01019173
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The
                                                                                                                    This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF0010-ABF9989, ABH0010-ABF9989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from MIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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  Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                       claim 1; SEQ ID NO 76290; 29pp + Sequence Listing; German.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genemic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                            Score 9.4; DB 1; Length 13; Pred. No. 1.3e+03; 0; Mismatches 1; Indels
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                                                                                                            Sequence 13 BP; 6 A; 4 C; 1 G; 2 T; 0 U; 0 Other;
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                                                                                                                                          Query Match
Best Local Similarity 90.9%;
Matches 10; Conservative
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 capeseen the oligomers described in the invantion. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                       SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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946 GGTTTAATGTATC 958
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                                1 GGTTTAATGGGTY
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21-FEB-2002 (first entry)

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12.9%; Score 9.4; DB 1; Length 13; llarity 76.9%; Pred. No. 1.3e+03; Conservative 1; Mismatches 2; Indels

Query Match Best Local Similarity Matches 10; Conserv

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99899, ABF00010-ABF99899, ABH00010-ABH99989 and ABI00010-ABI82073 tepresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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designed to detect single-nucleotide polymorphisms and cytosine
methylation status.
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Pred. No. 1.3e+03;
1; Mismatches 2; Indels
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ID ABF03754 standard; DNA; 13 BP.
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Best Local Similarity 76.9
Matches 10; Conservative
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                                                                      (EPIG-) EPIGENOMICS
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                                           SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Oligonucleotide SEQ ID NO 78050 for detecting SNP TSC0019867.
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Best Local Similarity
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                                                                        This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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Claim 1; SEQ ID NO 103751; 29pp + Sequence Listing; German
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Pred. No. 1.3e+03;
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Sequence 13 BP; 8 A; 3 C; 0 G; 2 T; 0 U; 0 Other;
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Matches 10; Conservative
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                                                                                                                 SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                            Oligonucleotide SEQ ID NO 5633 for detecting SNP TSC0001852,
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, contrant are also used for detecting cell type differentiation. ABC00010 aABC99989, ABF00010-ABF9989, ABH00010-ABH9989 and ABI00010-ABI82073 the represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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Matches 10; Conservative
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                                                                                                                                                                                                                                                                                       This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF99899, ABH00010-ABH99899 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPD at ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                 onucleotides, useful for diagnosis and cell typing, is detect single-nucleotide polymorphisms and cytosine
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Pred. No. 1.3e+03;
1; Mismatches 2; Indels
                                                                                                                                                                                                                                     Claim 1; SEQ ID NO 30480; 29pp + Sequence Listing; German.
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Best Local Similarity 76.9
Matches 10; Conservative
                                                                                                                       Set of oligonucleotides,
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                                                                                                                                                      designed to
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               range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99889, ABF00010-ABF99889, ABH00010-ABH99888 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electroic format from WIPO at fig. Mipo int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SNP, single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
oligonuclectides are used for diagnosis and/or prognosis of cancer and
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12.9%; Score 9.4; DB 1; Length 13;

Query Match

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI32073 tepsesent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at
                                                                                                                               SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                   Oligonucleotide SEQ ID NO 58153 for detecting SNP TSC0015616.
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Best Local Similarity 90.9
Matches 10; Conservative
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ABC84254
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Pred. No. 1.3e+03;
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                  Best Local Similarity 90.9%;
Matches 10; Conservative
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12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03; ive 0; Mismatches 1; Indels
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytoshie methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010 ABC099989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073
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                                                                                                                                                                                                                                This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99899, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic form part of the printed specification, but ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                         onucleotides, useful for diagnosis and cell typing, i detect single-nucleotide polymorphisms and cytosine
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                                                                                                                                                                                                         Claim 1; SEQ ID NO 84271; 29pp + Sequence Listing; German.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, contral nervous system, cardiovascular and metabolic disorders. The coligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF0010-ABF9989, ABH0010-ABH9989 and ABI0010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                        Claim 1; SEQ ID NO 84272; 29pp + Sequence Listing; German.
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90.9%;
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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12.9%; Score 9.4; DB 1; Length 13;
Best Local Similarity 90.9%; Pred. No. 1.3e+03;
Matches 10; Conservative 0; Mismatches 1; Indels
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12.9%; Score 9.4; DB 1; Length 13;
Best Local Similarity 90.9%; Pred. No. 1.3e+03;
Matches 10; Conservative 0; Mismatches 1; Indels
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AC ABC8823
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Berlin K;

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF9989, ABH00010-ABH9989 and ABI00010-ABI82073 topsement the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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Matches 10; Conservative
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genemic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC09989, ABF00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
SNP, single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99889, ABF00010-ABF99889, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; Ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                         set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                                                          Claim 1, SEQ ID NO 39918; 29pp + Sequence Listing; German.
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                                             Berlin K;
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            (EPIG-) EPIGENOMICS AG
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Sequence 13 BP; 2 A; 1 C; 4 G; 6 T; 0 U; 0 Other;

schultz1-899.rng

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABH99899 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo.int/pub/published_pct_sequences
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Sequence 13 BP; 4 A; 1 C; 4 G; 4 T; 0 U; 0 Other;

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12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03; ive 0; Mismatches 1; Indels
                               10; Conservative
                                                             955 TATCGCTACCA 965
                                                                                        11 TATCGCTAACA 1
Query Match
Best Local Similarity
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Gaps

RESULT 1868

ABF19666 standard; DNA; 13 BP. ABF19666;

(first entry) 21-FEB-2002 Oligonucleotide SEQ ID NO 119663 for detecting SNP TSC0029865.

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Homo sapiens.

WO200177384-A2

18-OCT-2001.

06-APR-2001; 2001WO-IB000713.

07-APR-2000; 2000DE-01019173.

(EPIG-) EPIGENOMICS AG

Berlin K; Olek A, Piepenbrock C,

WPI; 2001-657177/75.

Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.

Claim 1; SEQ ID NO 119663; 29pp + Sequence Listing; German.

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99889, ABF00010-ABF99899, ABH0010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence was obtained in electronic format from WIPD at the printed specification, but ftp.wipo.int/pub/published_pct_sequences ABP19666

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designed to detect single-nucleotide polymorphisms and cytosine
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          12.9%; Score 9.4; DB 1; Length 13; larity 90.9%; Pred. No. 1.38+03; . Conservative 0; Mismatches 1; Indels
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                                                                                                                                    ABF24055 standard; DNA; 13 BP.
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Query Match
Best Local Similarity
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Gaps ; 12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03; ive 0; Mismatches 1; Indels Sequence 13 BP; 2 A; 4 C; 0 G; 7 T; 0 U; 0 Other; Query Match
12.9
Best Local Similarity 90.9
Matches 10; Conservative 918 TCTTTGCCTTT 928 1 TCTTTACCTTT 11 g: 8

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic discorders. The oligomers are also used for detecting cell type differentiation. ABC0010 ABC099889, ABF00010-ABF99899 and ABI00010-ABF32073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at

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RESULT 1870 ABF31384

WO200177384-A2

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, aradiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABC0010-ABF9989, ABH0010-ABH99999 and ABT00010-ABI82073 trepresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from MIPO at
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                                                                                                                                                                                                                   SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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ABF31384 standard; DNA; 13
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The coligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99999, ABF00010-ABF99999, ABF00010-ABF999999, ABF00010-ABF999999, ABF00010-ABF99999, ABF00010-ABF999999, ABF00010-ABF99999, ABF000010-ABF99999, ABF000010-ABF99999, 
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                                                                                                                                    This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretracted genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABH99989 and ABI00010-ABI82073 tepresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but they wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                        Claim 1; SEQ ID NO 135478; 29pp + Sequence Listing; German.
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WPI; 2001-657177/75.
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central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99899. ABF00010-ABF9989. ABF00010-ABF9989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at fine bublished_pct_sequences.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99999, ABF00010-ABF99999, ABH00010-ABH99989 and ABI00010-ABI32073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                               SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                                                                                                                                 Oligonucleotide SEQ ID NO 219148 for detecting SNP TSC0053288.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99999, ABF00010-ABF99999, ABH00010-ABH99999 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but they wispo.int/pub/published_pct_sequences
                                         SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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designed to detect single-nucleotide polymorphisms and cytosine
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           Oligonucleotide SEQ ID NO 170119 for detecting SNP TSC0042467.
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ABF71267 standard; DNA; 13 BP.
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                                                                                                               (EPIG-) EPIGENOMICS AG
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010 ABC0999, ABC00010-ABC99989, ABC00010-ABC99989, ABC00010-ABC9989, ABC0010-ABC9989, ABC9989, ABC9
                                                          This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF0010-ABF9989, ABH0010-ABF9989 and ABI00010-ABI82073 tepseent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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designed to detect single-nucleotide polymorphisms and cytosine
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Claim 1; SEQ ID NO 171264; 29pp + Sequence Listing; German.
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ftp.wipo.int/pub/published_pct_sequences
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Matches 10; Conservative
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                     Oligonucleotide SEQ ID NO 175621 for detecting SNP TSC0043631.
                                       ABF75624 standard; DNA; 13 BP.
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                                                                                                                                                                                                                                                                                                                         SNP, single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                                   Oligonucleotide SEQ ID NO 200134 for detecting SNP TSC0049243.
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                                                              Score 9.4; DB 1; Length 13; Pred. No. 1.3e+03; 0; Mismatches 1; Indels
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                                       Seguence 13 BP; 4 A; 0 C; 4 G; 5 T; 0 U; 0 Other;
was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                    ABH00157 standard; DNA; 13 BP.
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90.9%;
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                                                                                             10; Conservative
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Best Local Similarity
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designed to detect single-nucleotide polymorphisms and cytosine
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                                                                                                                                                                                                                                                                                      Claim 1; SEQ ID NO 175621; 29pp + Sequence Listing; German.
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Pred. No. 1.3e+03;
1; Mismatches 2; Indels
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                                                     Berlin K;
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76.9%;
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Best Local Similarity 76.9
Matches 10, Conservative
                                                     Piepenbrock C,
(EPIG-) EPIGENOMICS AG
                                                                                                                WPI; 2001-657177/75
                                                                                                                                                                                                                                    methylation status.
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Score 9.4; DB 1; Length 13; Pred. No. 1.3e+03; 0; Mismatches 1; Indels

12.9%; 90.9%;

Query Match 12.9 Best Local Similarity 90.9 Matches 10; Conservative

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                              set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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designed to detect single-nucleotide polymorphisms and cytosine
methylation status.
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                                                                                                                                                                                         Claim 1; SEQ ID NO 203371; 29pp + Sequence Listing; German.
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                  Berlin K;
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12.9%;
Best Local Similarity 90.9%;
Matches 10; Conservative
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                    Piepenbrock
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central nervous system; gastrointestinal; respiratory; immune; metabolic.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, contral nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but fip.wipo.int/pub/published_pct_sequences
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Score 9.4; DB 1; Length 13;
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             oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligoners are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF9989, ABH00010-ABF99989 and ABI00010-ABF82073 represent the oligoners described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but they was obtained in electronic format from WIPO at
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ABH35003 standard; DNA; 13 BP.

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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC09989, ABC0010-ABE99899, ABC0010-ABE99989, ABC0010-ABE99989, ABC0010-ABE99989, and ABI00010-ABE82073 represent the oligomers described in the invention. NOTE: The sequence was obtained in electronic form part of the printed specification, but the wipo.int/pub/published_pct_sequences
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                                                                                                                         SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                        Oligonucleotide SEQ ID NO 234980 for detecting SNP TSC0057373.
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-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at

Page 836

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Gaps

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12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03; ive 0; Mismatches 1; Indels

10; Conservative

Matches

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Query Match

947 GTTTAATGTAT 957

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Sequence 13 BP; 5 A; 4 C; 0 G; 3 T; 0 U; 1 Other;

was obtained in electronic format from Wi ftp.wipo.int/pub/published_pct_sequences

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Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine
                                                                     Local Similarity
                                                                                                                                                                                                   methylation status.
                                                                                                                                                 WO200177384-A2
                                                                                                                                           Homo sapiens
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                                                                                                           ABF60977;
                                                                  Query Match
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                                                                                              Matches
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nuclectide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABE99899, ABF00010-ABE99899, ABF00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo.int/pub/published_pct_sequences
onucleotides, useful for diagnosis and cell typing, i
detect single-nucleotide polymorphisms and cytosine
                                                                                                                                                                                         German.
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                                                                                                                                                                                         Claim 1; SEQ ID NO 236088; 29pp + Sequence Listing;
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oligonucleotides, useful
                                        designed to detect methylation status.
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                                                                                           ABF60977 standard; DNA; 13
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Oligonucleotide SEQ ID NO 160974 for detecting SNP TSC0005250. 22-FEB-2002 (first entry)

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

06-APR-2001; 2001WO-IB000713

07-APR-2000; 2000DE-01019173

(EPIG-) EPIGENOMICS AG

Berlin K; Piepenbrock C, WPI; 2001-657177/75. Claim 1; SEQ ID NO 160974; 29pp + Sequence Listing; German.

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligoners for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastroinfestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010

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Gaps

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948 TITAAIGIAICGC 960

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Local Similarity 76.9 nes 10; Conservative

Matches

·Query Match

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABF32073 represent the oligomers described in the invention. NOTE: The sequence data for this patent din out form part of the printed specification, but the wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                         SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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designed to detect single-nucleotide polymorphisms and cytosine
                                                                                                                                                                                                      Oligonucleotide SEQ ID NO 161729 for detecting SNP TSC0040712.
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                                                                                             ABF61732 standard; DNA; 13
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting call type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABT00010-ABI82073 trepresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
              SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                                                                                                                                                                                                         SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                               Oligonucleotide SEQ ID NO 212091 for detecting SNP TSC0051687.
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12.9%; Score 9.4; DB 1; Length 13;
Best Local Similarity 76.9%; Pred. No. 1.3e+03;
Matches 10; Conservative 1; Mismatches 2; Indels
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1 ATTGGTGTTATGY 13
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but fup.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                     Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                Berlin K;
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          This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99999, ABF00010-ABF99999, ABH00010-ABH99999 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic formmat from WIPO at
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                                            Score 9.4; DB 1; Length 13;
Pred. No. 1.3e+03;
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RESULT 1897

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF0010-ABF9989, ABH0010-ABH9989 and ABI00010-ABI82073 trepresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from NIPO at
                                                                                                                                                                    SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                                              Oligonucleotide SEQ ID NO 253738 for detecting SNP TSC0061857.
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BP.
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ABH53761 standard; DNA; 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                       Score 9.4; DB 1; Length 13
Pred. No. 1.3e+03;
0; Mismatches 1; Indels
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BP; 3 A; 2 C; 0 G; 8 T; 0 U; 0 Other;
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                                          Query Match
Best Local Similarity 90.9%;
Matches 10; Conservative
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Matches 10; Conserv
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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Matches 10, Conservative
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Pred. No. 1.38+03;
0; Mismatches 1; Indels
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  Homo sapiens.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF9989, ABH00010-ABH9989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                    SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                      Oligonucleotide SEQ ID NO 42402 for detecting SNP TSC0012648.
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central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABR00010-ABR99989, ABR00010-ABR99989, ABR00010-ABR99989 and ABR00010-ABR99989 are represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; 88; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                        Oligonucleotide SEQ ID NO 93048 for detecting SNP TSC0023263
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                                                                                This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nuclectide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010 -ABC99989, ABF0010-ABF99899, ABH00010-ABH99899 and ABI00010-ABIS2073 data for this patent did not form part of the printed specification, but ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03; ive 0; Mismatches 1; Indels
                                          Claim 1; SEQ ID NO 49359; 29pp + Sequence Listing; German.
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methylation status.
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data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                 12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03;
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ABF02725 standard; DNA; 13
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                                                                                                                                          SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                     Oligonucleotide SEQ ID NO 52802 for detecting SNP TSC0014620.
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ABC52785 standard; DNA; 13 BP
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peptide nucleic acid, cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretracted genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, acadiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989 in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but fitp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                            Claim 1; SEQ ID NO 54062; 29pp + Sequence Listing; German.
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Matches 10; Conservative
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(EPIG-) EPIGENOMICS
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acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrolinestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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designed to detect single-nucleotide polymorphisms and cytosine
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                                                                                                                                                                                                                                             Sequence 13 BP; 8 A; 4 C; 0 G; 0 T; 0 U; 1 Other;
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and oytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99889, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                          SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                         Oligonuclectide SEQ ID NO 32325 for detecting SNP TSC0010079.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; 88; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                       Score 9.4; DB 1; Length 13; Pred. No. 1.3e+03; 0; Mismatches 1; Indels
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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1; Indels
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ABC32308 standard; DNA; 13 BP

RESULT 1915 ABC32308 ID ABC32301

Score 9.4; DB 1; Length 13; Pred. No. 1.36+03;

12.98;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Pred. No. 1.3e+03;
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Best Local Similarity 90.9
Matches 10; Conservative
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF99899, ABH00010-ABF99899 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic formmat from WIPO at the printed specification, but ftp.wipo.int/pub/published_pct_sequences
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Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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Pred. No. 1.3e+03;
0; Mismatches 1; Indels
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                                                                                                                                                                               Claim 1; SEQ ID NO 83394; 29pp + Sequence Listing; German.
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Matches 10; Conservative
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oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF0010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NoTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at flow they bublished_pct_sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SNP, single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                      SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Query Match 12.9%; Score 9.4; DB 1; Length 13; Best Local Similarity 90.9%; Pred. No. 1.3e+03; Matches 10; Conservative 0; Mismatches 1; Indels

Piepenbrock C,

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Homo sapiens

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This invention describes novel oligonucleotide primers or peptide nucleic
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designed to detect single-nucleotide polymorphisms and cytosine
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Matches 10; Conservative
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                              SNP; single nuclectide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Oligonucleotide SEQ ID NO 62962 for detecting SNP TSC0016655.
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Pred. No. 1.3e+03;
0; Mismatches 1; Indels
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Homo sapiens

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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Claim 1; SEQ ID NO 15502; 29pp + Sequence Listing; German.
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                                                            Score 9.4; DB 1; Length 13;
Pred. No. 1.3e+03;
1; Mismatches 2; Indels
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                                                                                                                 SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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designed to detect single-nucleotide polymorphisms and cytosine
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                                                                                            Oligonucleotide SEQ ID NO 137355 for detecting SNP TSC0034314.
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                     ABF37358 standard; DNA; 13
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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Berlin

Piepenbrock C,

Olek A,

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligoners for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligoners are also used for detecting cell type differentiation. ABC00010-ABC99989, ABR00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                       oligonucleotides, useful for diagnosis and cell typing, is to detect single-nucleotide polymorphisms and cytosine
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99889, ABF00010-ABF99899, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                               SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                          Oligonucleotide SEQ ID NO 149160 for detecting SNP TSC0037626.
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                                                                                                                                                                                                                                                                                                                                                                                           SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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  90.9%; Pred. No. 1.3e+03; ive 0; Mismatches 1; Indels
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acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory. Central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99389, ABF0010-ABF99989, ABH0010-ABH99989 and ABI00010-ABI82073. Tepresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at the printed specification, but fire wipo.int/pub/published_pct_sequences
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Pred. No. 1.3e+03;
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designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                              Claim 1; SEQ ID NO 227267; 29pp + Sequence Listing; German.
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SND; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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designed to detect single-nucleotide polymorphisms and cytosine
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                                                                        ABH04878 standard; DNA; 13 BP.
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represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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Matches 10; Conservative 0; Mismatches 1; Indels
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Berlin K;

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABF99989 and ABI00010-ABF92073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic formmat from WIPO at fitte.wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 13 BP; 1 A; 0 C; 4 G; 8 T; 0 U; 0 Other;
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Matches 10; Conservative
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Local Similarity 90.9 Les 10; Conservative 913 TTTGGTCTTTG 923

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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                                                                                                                                         Claim 1; SEQ ID NO 208262; 29pp + Sequence Listing; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03; tive 0; Mismatches 1; Indels
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Matches 10, Conservative
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                                                                                                                                                                                                                                                                                                     WPI; 2001-657177/75
                                                                                                                                                                                                                                                                                                                                                       designed to detect methylation status.
                                                                                                  WO200177384-A2
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at fitp.wipo.int/pub/published_pot_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               12.9%; Score 9.4; DB 1; Length 13; 76.9%; Pred. No. 1.3e+03;
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              (EPIG-) EPIGENOMICS AG.
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Claim 1; SEQ ID NO 209246; 29pp + Sequence Listing; German.

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Sequence 13 BP; 1 A; 1 C; 4 G; 7 T; 0 U; 0 Other;

Sequence 13 BP; 2 A; 4 C; 0 G; 7 T; 0 U; 0 Other;

0; Gaps 12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03; ive 0; Mismatches 1; Indels 0uery Match Best Local Similarity 90.9 Matches 10, Conservative

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ABH09784 standard; DNA; 13 BP. RESULT 1941 ABH09784

ABH09784;

(first entry) 22-FEB-2002 Oligonuclectide SEQ ID NO 209761 for detecting SNP TSC0051215.

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Homo sapiens.

WO200177384-A2.

18-OCT-2001.

06-APR-2001; 2001WO-IB000713.

07-APR-2000; 2000DE-01019173.

(EPIG-) EPIGENOMICS AG.

Berlin Olek A, Piepenbrock C,

WPI; 2001-657177/75.

Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.

Claim 1; SEQ ID NO 209761; 29pp + Sequence Listing; German.

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99899, ABF00010-ABF99899, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at

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                                                                                                                                               SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                 Gaps
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                                                                                                                                Oligonucleotide SEQ ID NO 237059 for detecting SNP TSC0057828.
Score 9.4; DB 1; Length 13;
Pred. No. 1.3e+03;
0; Mismatches 1; Indels
                                                                                     ABH37082 standard; DNA; 13 BP.
 12.9%;
90.9%;
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Query Match
Best Local Similarity 90.9
Matches 10; Conservative
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ABH37082/c
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Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine WPI; 2001-657177/75. methylation status.

Berlin K;

Olek A, Piepenbrock C,

(EPIG-) EPIGENOMICS AG.

06-APR-2001; 2001WO-IB000713. 07-APR-2000; 2000DE-01019173

WO200177384-A2

18-OCT-2001.

Claim 1; SEQ ID NO 237059; 29pp + Sequence Listing; German.

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically prerzeated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC09989, ABC0010-ABF99899, ABH0010-ABH99999 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic formmat from WIPO at fire.wipo.int/pub/published_pct_sequences

Sequence 13 BP; 2 A; 1 C; 4 G; 6 T; 0 U; 0 Other;

Gaps ; 0 Match 12.9%; Score 9.4; DB 1; Length 13; Local Similarity 90.9%; Pred. No. 1.3e+03; les 10; Conservative 0; Mismatches 1; Indels Query Match Matches

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960 CIACCAACGGT 970 CTACCAACGAT 2 12 qq

RESULT 1943 ABF62508

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ABF62508 standard; DNA; 13

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                                                                                                     SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                  Oligonucleotide SEQ ID NO 162505 for detecting SNP TSC0040879.
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ABH38409 standard; DNA; 13
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  22-FEB-2002
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The coligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 the represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 13 BP; 8 A; 2 C; 0 G; 2 T; 0 U; 1 Other;
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                                                                                                                                                                              Piepenbrock C,
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les 10; Conservative
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                                                                                                                                                                           This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABH32073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting call type differentiation. ABC0010-ABC39989, ABF00010-ABF99989, ABF00010-ABF9989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at the printed specification, but ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                                                                                                               SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Pred. No. 1.3e+03;
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                                                                    SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Oligonucleotide SEQ ID NO 248714 for detecting SNP TSC0060779.
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This invention describes novel oligonucleotide primers or peptide mucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genemic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99899, ABF00010-ABF99989 and metabolic disorders. The represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo.int/pub/published_pct_sequences
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                                                                                                                                                             onucleotides, useful for diagnosis and cell typing, is detect single-nucleotide polymorphisms and cytosine
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                                  07-APR-2000; 2000DE-01019173
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Best Local Similarity 76.9
Matches 10; Conservative
                                                                                                                                                              Set of oligonucleotides,
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                                                                                             Olek A, Piepenbrock C,
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00110-ABC99999, ABF00010-ABF99999, ABH00010-ABH99999 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic formmat from WIPO at
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                             Claim 1; SEQ ID NO 92858; 29pp + Sequence Listing; German.
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Best Local Similarity
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                    Sequence 13 BP; 1 A; 0 C; 4 G; 7 T; 0 U; 1 Other;
was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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76.9%;
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                                                   Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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acid (PNA) oligomers for detecting single nuclectide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99889, ABC0010-ABE9988, ABC0010-ABE9988, ABC0010-ABE9988, and ABE0010-ABE82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                                                                                                                                                         This invention describes novel oligonucleotide primers or peptide nucleic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                              Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine
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                                                                                                                                                                                                          Claim 1; SEQ ID NO 50417; 29pp + Sequence Listing; German.
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                                                                                                                                                             methylation status.
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Pred. No. 1.3e+03;
0; Mismatches 1; Indels
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABC9989, ABF00010-ABG9989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but they was obtained in electronic format from WIPO at
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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ftp.wipo.int/pub/published_pct_sequences
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BP.

ABF02724 standard; DNA; 13

ABF02724 ID ABF0 XX

Sequence 13 BP; 2 A; 0 C; 3 G; 7 T; 0 U; 1 Other;

RESULT 1960

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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99889, ABF00010-ABF99889, ABF00010-ABF99899, ABF00010-ABF99899 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at the printed specification, but ftp.wipo.int/pub/published_pot_sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Oligonucleotide SEQ ID NO 28626 for detecting SNP TSC0008250.
                                                                                                                                                                                                                                                                                                                                                                                    Claim 1; SEQ ID NO 3274; 29pp + Sequence Listing; German.
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Pred. No. 1.3e+03;
0; Mismatches 1;
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ilarity 90.9%;
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                                                                                                                                 SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                        Oligonucleotide SEQ ID NO 102721 for detecting SNP TSC0025656.
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90.9%;
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Matches 10; Conservative
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acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                  Claim 1; SEQ ID NO 28626; 29pp + Sequence Listing; German.
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-ABC99989, ABF00010-ABF99989, ABH00010-ABH99889 and ABI00010-ABI82073 represent the oligomers described in the invention. NoTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                 Seguence 13 BP; 7 A; 5 C; 0 G; 0 T; 0 U; 1 Other;
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastronitestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010

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                                                                                                                                                               SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, coincomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF99889, ABH0010-ABH9989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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         (EPIG-) EPIGENOMICS AG
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Claim 1; SEQ ID NO 83299; 29pp + Sequence Listing; German.

Berlin K;

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABH00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                             Score 9.4; DB 1; Length 13; Pred. No. 1.3e+03; 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                     ABC14556 standard; DNA; 13 BP
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acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and range of diseases including immune system, gastrointestinal, respiratory. Central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC099889, ABC0010-ABC99889 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at the printed specification, but fire.wipo.int/pub/published_pct_sequence
                                                                                                                                          This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP)
                                             Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                            Claim 1; SEQ ID NO 90624; 29pp + Sequence Listing; German.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a
                                                                                          SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Oligonucleotide SEQ ID NO 41073 for detecting SNP TSC0012383.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The coligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99999, ABF00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at fire.wipo.int/pub/published_pct_sequences
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range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC099889, ABF00010-ABF99889, ABF00010-ABH99889 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form par of the printed specification, but was obtained in electronic format from WIPO at the printed specification, but ftp.wipo.int/pub/published_pct_sequences
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Pred. No. 1.3e+03;
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                                                                                                                                                                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; 88; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                        SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                    Oligonucleotide SEQ ID NO 128319 for detecting SNP TSC0032146.
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Homo sapiens

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic discorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABE99989, ABF00010-ABE99989, ABF00010-ABE99989, ABF00010-ABE99989, abstroint cell type differentiation. ABC00010 data for this patent did not form part of the printed specification, but was obtained in electronic formmat from WIPO at
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                                                                                                                                                                                                                                                                                    Claim 1; SEQ ID NO 131783; 29pp + Sequence Listing; German.
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                                                   This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABE99989, ABF00010-ABE99989, ABF00010-ABE99989, ABF00010-ABE99989, and ABI00010-ABE9073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                            Claim 1; SEQ ID NO 133099; 29pp + Sequence Listing; German.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, contral nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99889, ABH00010-ABH99989 and ABI00010-ABI82073 tepseson the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                   SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                             ABF71904 standard; DNA; 13
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nes 10; Conservative
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RESULT 1982
ABP71904
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ABP71904
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ABP71900
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DE Oligomu
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SNP, sin
WC2017
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HOMO SE
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99889, ABF00010-ABF99899 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but twipo.int/pub/published_pct_sequences
peptide nucleic acid, cytosine methylation; cardiovascular; primer; ss;
central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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RESULT 1986
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                                                                                                                                  This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) eligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The eligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The eligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                       set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                               Claim 1; SEQ ID NO 175622; 29pp + Sequence Listing; German.
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                       Olek A, Piepenbrock C,
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABH00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from MIPO at
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acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immine system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC09989, ABC0010-ABF9989, ABH0010-ABH99989 and ABI0010-ABI82073. represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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Pred. No. 1.3e+03;
0; Mismatches 1;
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(first entry)

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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                             Oligonucleotide SEQ ID NO 180149 for detecting SNP TSC0044601.
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                      Length 13;
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                      12.9%;
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF99989, ABH00010-ABF99989 and ABI00010-ABF3073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic formmat from WIPO at the printed specification, but fire wipo.int/pub/published_pot_sequences
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Pred. No. 1.3e+03;
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically precreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic discorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine methylation status.
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oligomers are also used for detecting cell type differentiation. ABC00010 -ABC9989, ABF0010-ABF99999, ABF00010-ABF99999, ABF0010-ABF99999 and ABI00010-ABF92073 represent the oligomers described in the invention. NoTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at frow.wipo.int/pub/published_pct_sequences
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                        SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                  SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Oligonucleotide SEQ ID NO 235462 for detecting SNP TSC0057483.
                                                                                                                                                                                                                                                                                                        Claim 1; SEQ ID NO 235462; 29pp + Sequence Listing; German.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABE99989, ABF00010-ABE99989, ABH00010-ABE99989 and ABI00010-ABE3073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo.int/pub/published_pct_sequences
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Claim 1; SEQ ID NO 212322; 29pp + Sequence Listing; German.
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                                 12.9%; Score 9.4; DB 1; Length 13; llarity 90.9%; Pred. No. 1.3e+03; Conservative 0; Mismatches 1; Indels
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from NIPO at
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic discorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF99989, ABH00010-ABF99989, and ABI00010-ABF9073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but two botained in electronic format from WIPO at
oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NoTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at flow int/pub/published_pot_sequences
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                                                                                                                                                                                                                             12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03; tive 0; Mismatches 1; Indels
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Pred. No. 1.3e+03;
1; Mismatches 2; Indels
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12.9%; Score 9.4; DB 1; Length 13;

Query Match

Sequence 13 BP; 4 A; 4 C; 0 G; 4 T; 0 U; 1 Other;

ABH65132,

à g

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF99989, ABH00010-ABF99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic formmat from WIPO at fitted specification, but fit wipo.int/pub/published_pct_sequences
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                                                                                                   SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                              Oligonucleotide SEQ ID NO 42546 for detecting SNP TSC0012678.
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                     21-FEB-2002 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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76.9%; Pred. No. 1.3e+03; indels ive 1; Mismatches 2; Indels
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designed to detect single-nucleotide polymorphisms and cytosine
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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designed to detect single-nucleotide polymorphisms and cytosine
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                                                                                                                                  Claim 1; SEQ ID NO 22224; 29pp + Sequence Listing; German
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represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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Matches 10; Conservative 0; Mismatches 1; Indels
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                                                                                                 Sequence 13 BP; 9 A; 0 C; 1 G; 3 T; 0 U; 0 Other;
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                         SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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ABC54212 standard; DNA; 13
                             ABF01571 standard; DNA; 13
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Claim 1; SEQ ID NO 105981; 29pp + Sequence Listing; German.
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, asrdiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99899, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at
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                                                                      Berlin K;
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Length 13;

G; 6 T; 0 U; 0 Other;

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A; 5

Sequence 13 BP; 2

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABR99899, ABR00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genemic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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designed to detect single-nucleotide polymorphisms and cytosine
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/ Match 12.9%; Score 9.4; DB 1; Length 13 Local Similarity 90.9%; Pred. No. 1.3e+03; nes 10; Conservative 0; Mismatches 1; Indels
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methylation status.
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RESULT 2016

ABC86604

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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                               Oligonucleotide SEQ ID NO 86621 for detecting SNP TSC0021768.
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ftp.wipo.int/pub/published_pct_sequences
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ABC86604 standard; DNA; 13
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                     ABC86604
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genemic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010 +ABC99989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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Gaps
Score 9.4; DB 1; Length 13;
Pred. No. 1.3e+03;
1; Mismatches 2; Indels
                                                                         946 GGTTTAATGTATC 958
Query Match 12.9
Best Local Similarity 76.9
Matches 10; Conservative
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ABC12758 standard; DNA; 13 (first entry) 1 GGTTTAAAGGATY 13 20-FEB-2002 ABC12758 ABC12758
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Homo sapiens

Oligonucleotide SEQ ID NO 12765 for detecting SNP TSC0002990.

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99899, ABF00010-ABF99989, ABH00010-ABH99999 and ABI00010-ABF8073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo int/pub/published_pct_sequences Gaps set or oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status. ö Score 9.4; DB 1; Length 13; Pred. No. 1.3e+03; 1; Mismatches 2; Indels Claim 1; SEQ ID NO 12765; 29pp + Sequence Listing; German. Sequence 13 BP; 0 A; 0 C; 2 G; 10 T; 0 U; 1 Other; Berlin K; 06-APR-2001; 2001WO-IB000713. 12.9%; 07-APR-2000; 2000DE-01019173 TICTITGGICITI 922 Local Similarity 76.9 nes 10; Conservative Olek A, Piepenbrock C, (EPIG-) EPIGENOMICS AG WPI; 2001-657177/75 WO200177384-A2. 18-OCT-2001 910 Query Match RESULT 2018 Matches ABC63662 ઠ g

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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic. Oligonucleotide SEQ ID NO 63679 for detecting SNP TSC0016816. WO200177384-A2. Homo sapiens

BP.

ABC63662 standard; DNA; 13

(first entry)

21-FEB-2002

ABC63662;

Olek A, Piepenbrock C, Berlin K; 07-APR-2000; 2000DE-01019173. (EPIG-) EPIGENOMICS AG

06-APR-2001; 2001WO-IB000713.

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                                                                                            This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                       Claim 1; SEQ ID NO 63679; 29pp + Sequence Listing; German.
                        tides, useful for diagnosis and cell single-nucleotide polymorphisms and
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABE99989, ABF00010-ABE99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABP9989 and ABIO0010-ABP192073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at fire wipo.int/pub/published_pct_sequences
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Best Local Similarity 90.9
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                                                                                                                                                                                                                                          SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABH9989 and ABI00010-ABI82073 trapseent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from MIPO at
                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                             Oligonucleotide SEQ ID NO 42132 for detecting SNP TSC0012592.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABF09999, ABF00010-ABF9999, ABH00010-ABF9999, ad ABI00010-ABF9999, abround are described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but
                                             This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, oardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABE99989, ABF00010-ABE9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                Claim 1; SEQ ID NO 133389; 29pp + Sequence Listing; German.
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                                                                                                                                                                                                                                                                      Claim 1; SEQ ID NO 117076; 29pp + Sequence Listing; German.
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ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                      12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03;
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic formmat from WIPO-at
                                                                                                                                                                                                                          SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and oycosite methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99889, ABF00010-ABF99899, ABH00010-ABF99989, ABH00010-ABF99989 and ABI00010-ABF3073 data for this patent did not form part of the printed specification, but was obtained in electronic formmat from WIPO at
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central nervous system; gastrointestinal; respiratory; immune; metabolic.
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and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF9989, ABH00010-ABH99999 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                              SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                        Oligonucleotide SEQ ID NO 226937 for detecting SNP TSC0055323.
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12.9%; Score 9.4; DB 1; Length 13;
Best Local Similarity 90.9%; Pred. No. 1.3e+03;
Matches 10; Conservative 0; Mismatches 1; Indels
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  ABH26960;
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, certral nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99889, ABF00010-ABF99889, ABF00010-ABF99889 and ABI00010-ABF8073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                Claim 1; SEQ ID NO 227327; 29pp + Sequence Listing; German.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                  represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at the printed specification, the ftp.wipo.int/pub/published_pct_sequences
  -ABC99988, ABF00010-ABF99988, ABH00010-ABH99988 and ABI00010-ABI82073
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Pred. No. 1.3e+03;
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90.9%;
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                                                                                                                                                          SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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12.9%; Score 9.4; DB 1; Length 13;
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ftp.wipo.int/pub/published_pct_sequences
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                                                                  ABF53194 standard; DNA; 13
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF0010-ABF99989, ABH0010-ABF99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Matches 10; Conservative
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                           Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                             Berlin K;
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07-APR-2000; 2000DE-01019173
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                                                             Piepenbrock C,
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Matches 10; Conserv
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Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.

Berlin K;

Piepenbrock C,

olek A,

WPI; 2001-657177/75

(EPIG-) EPIGENOMICS AG

06-APR-2001; 2001WO-IB000713. 07-APR-2000; 2000DE-01019173

18-OCT-2001,

Claim 1; SEQ ID NO 205296; 29pp + Sequence Listing; German.

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           This invention describes novel oligonucleotide primars or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99899, ABF00010-ABF99899, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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Pred. No. 1.3e+03;
0; Mismatches 1; Indels
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Best Local Similarity 90.9%;
Matches 10; Conservative
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(first entry)

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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; 88; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                  Oligonucleotide SEQ ID NO 186649 for detecting SNP TSC0045992.
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            ABF86652 standard; DNA; 13
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                                                                                                                                                                                                                                                                                                                                  SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; 88; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                         Query Match 12.9%; Score 9.4; DB 1; Length 13; Best Local Similarity 90.9%; Pred. No. 1.3e+03; Matches 10; Conservative 0; Mismatches 1; Indels
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              Sequence 13 BP; 7 A; 0 C; 5 G; 1 T; 0 U; 0 Other;
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90.9%;
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Best Local Similarity 90.9
Matches 10, Conservative
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Berlin K;

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and oytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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Claim 1; SEQ ID NO 186649; 29pp + Sequence Listing; German.
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Best Local Similarity 90.9
Matches 10; Conservative
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947 GTTTAATGTAT 957

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RESULT 2044

Homo sapiens.

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                                                                                                                                                                                                                                                                                                                                                                                             Claim 1; SEQ ID NO 212092; 29pp + Sequence Listing; German.
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WO200177384-A2
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting call type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic formmat from WIPO at
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                                              Set of oligonuclectides, useful for diagnosis and cell typing, is designed to detect single-nuclectide polymorphisms and cytosine methylation status.
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                                                                                                                         Claim 1; SEQ ID NO 162508; 29pp + Sequence Listing; German
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Pred. No. 1.3e+03;
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90.9%;
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC099889, ABF00010-ABF99889, ABF00010-ABH99889 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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Matches 10; Conservative
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Score 9.4; DB 1; Length 13; Pred. No. 1.3e+03;

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Query Match Best Local Similarity

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Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.

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Piepenbrock C,

olek A,

WPI; 2001-657177/75

(EPIG-) EPIGENOMICS

06-APR-2001; 2001WO-IB000713. 07-APR-2000; 2000DE-01019173. Claim 1; SEQ ID NO 191687; 29pp + Sequence Listing; German.

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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretraeted genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABH99989 and ABI00010-ABI82073 tepresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but they was obtained in electronic form at from WIPO at
                                                      SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                          Oligonucleotide SEQ ID NO 216146 for detecting SNP TSC0052566.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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Pred. No. 1.3e+03;
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

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Homo sapiens

Oligonucleotide SEQ ID NO 191687 for detecting SNP ISC0000813.

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                                                                           This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC099889, ABF00010-ABE99899, ABH00010-ABH99899 and ABI00010-ABE82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at the printed specification, but the wipo.int/pub/published_pct_sequences
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                                        Claim 1; SEQ ID NO 248597; 29pp + Sequence Listing; German.
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acid (PNA) oligomers for detecting single nucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomicleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABF00010-ABF9989, ABF00010-ABF9
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data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                      SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                 Oligonucleotide SEQ ID NO 263878 for detecting SNP TSC0063961.
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peptide nucleic acid, cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                       Olek A, Piepenbrock C,
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically prereated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, cointrain nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABR00010-ABF9989, ABH0010-ABH9989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABP9989, ABF00010-ABP9989, and ABI00010-ABB182073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretracted genomic DNA. The oligomucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF0010-ABF9989, ABH0010-ABH99989 and ABI00110-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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Score 9.4; DB 1; Length 13;
Pred. No. 1.3e+03;
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                               SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                 Oligonucleotide SEQ ID NO 97666 for detecting SNP TSC0024259.
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Pred. No. 1.3e+03;
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Best Local Similarity
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                                                                                                                                                                                   Homo sapiens
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               ABC97649;
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ABC97968/c
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ABC97649 standard; DNA; 13 BP.

RESULT 2061

ABC97649/ ID ABC9'

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Pred. No. 1.3e+03;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                         Claim 1; SEQ ID NO 97985; 29pp + Sequence Listing; German.
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                                                             06-APR-2001; 2001WO-IB000713
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Best Local Similarity 90.9
Matches 10; Conservative
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                                                                                                                                                     Piepenbrock C,
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WO200177384-A2
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                              18-OCT-2001
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                      Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                                                    Claim 1; SEQ ID NO 23969; 29pp + Sequence Listing; German
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Best Local Similarity 76.9
Matches 10; Conservative
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ABC23953/53/KX
ABC20-E7
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oligomers are also used for detecting cell type differentiation. ABC00010 -ABC9989, ABF00010-ABF99899, ABF00010-ABF99899, ABF00010-ABF99899, ABF00010-ABF99899, ABF00010-ABF99999 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at fixed in the printed specification, but ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, coingomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SNP; single nuclectide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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12.9%; Score 9.4; DB 1; Length 13;
Best Local Similarity 90.9%; Pred. No. 1.3e+03;
Matches 10; Conservative 0; Mismatches 1; Indels
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Matches 10; Conservative
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                                                                                                                                                                                                                                                        SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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designed to detect single-nucleotide polymorphisms and cytosine
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Matches 10; Conservative
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schultz1-899.rng

Berlin K;

Piepenbrock C,

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC09989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                     Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                                                                                                                                                                                                                                                                                                         Claim 1; SEQ ID NO 101567; 29pp + Sequence Listing; German.
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                             07-APR-2000; 2000DE-01019173
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                                                                                       (EPIG-) EPIGENOMICS AG.
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                                                   SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Oligonuclectide SEQ ID NO 1564 for detecting SNP TSC0000566.
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12.9%; Score 9.4; DB 1; Length 13;
Best Local Similarity 90.9%; Pred. No. 1.3e+03;
Matches 10; Conservative 0; Mismatches 1; Indels
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                                                                                Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                Oligonucleotide SEQ ID NO 2891 for detecting SNP TSC0001129.
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                                                         20-FEB-2002 (first entry)
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SNP, single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

06-APR-2001; 2001WO-IB000713

WO200177384-A2

18-OCT-2001

Homo sapiens

Oligonucleotide SEQ ID NO 101567 for detecting SNP TSC0025295.

(first entry)

21-FEB-2002

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ABF01570 standard; DNA; 13

RESULT 2068 **ABF0157**

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                                         This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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designed to detect single-nucleotide polymorphisms and cytosine
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Claim 1; SEQ ID NO 2891; 29pp + Sequence Listing; German.
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Best Local Similarity 90.9
Matches 10; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Pred. No. 1.3e+03;
1; Mismatches 2; Indels
                                                      Sequence 13 BP; 2 A; 0 C; 2 G; 8 T; 0 U; 1 Other;
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76.9%;
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                                                                                                                             SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                    Oligonucleotide SEQ ID NO 105160 for detecting SNP TSC0026342.
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                           ABF05163 standard; DNA; 13
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RESULT 2073
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretraeted genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC09989, ABF00010-ABE99899, ABF00010-ABE99899, ABF00010-ABE99899 and ABI00010-ABE8073 represent the oligomers described in the invention. NOTE: The sequence was obtained in electronic format from WIPO at the printed specification, but the wipo.int/pub/published_pct_sequences
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Matches 10; Conservative
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                                                                                                                                                        Claim 1; SEQ ID NO 109989; 29pp + Sequence Listing; German.
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range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The coligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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12.9%; Score 9.4; DB 1; Length 13;

Query Match

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(first entry)
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                                                                                                                                                                                                                 SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Pred. No. 1.3e+03;
0; Mismatches 1; Indels
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ABC86605 standard; DNA; 13
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 Best Local Similarity 90.8
Matches 10; Conservative
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABE99989, ABF00010-ABE99989, ABF00010-ABE99989 and ABI00010-ABE32073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WFPO at fitte.wipo.int/pub/published_pot_sequences
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                                                             SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Oligonucleotide SEQ ID NO 62367 for detecting SNP TSC0016537.
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acid (PNA) oligomers for detecting single muclectide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99898 ABH00010-ABF99899 and ABI00010-ABF99980 in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                      This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                        nnucleotides, useful for diagnosis and cell typing, i detect single-nucleotide polymorphisms and cytosine
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                                                                                                                                                                                                                                                           Claim 1; SEQ ID NO 62961; 29pp + Sequence Listing; German.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010 ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073
                                                                                       This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99889, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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designed to detect single-nucleotide polymorphisms and cytosine
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Pred. No. 1.3e+03;
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                                                           Claim 1; SEQ ID NO 88247; 29pp + Sequence Listing; German.
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Best Local Similarity 90.9
Matches 10, Conservative
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                          methylation status.
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                     Oligonucleotide SEQ ID NO 134391 for detecting SNP TSC0033498.
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                                                                                     ABF34394 standard; DNA; 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                     SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at fit, wipo.int/pub/published_pct_sequences
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                                                                                                                 Query Match
12.9%; Score 9.4; DB 1; Length 13;
Best Local Similarity 90.9%; Pred. No. 1.3e+03;
Matches 10; Conservative 0; Mismatches 1; Indels
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABE99989, ABF00010-ABE99989, ABH00010-ABE99989 and ABI00010-ABE82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo.int/pub/published_pct_sequences
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12.9%; Score 9.4; DB 1; Length 13;
Best Local Similarity 90.9%; Pred. No. 1.3e+03;
Matches 10; Conservative 0; Mismatches 1; Indels
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; 8s; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99999, ABF00010-ABF99999, ABH00010-ABH99999 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                           oligonucleotides, useful for diagnosis and cell typing, is at to detect single-nucleotide polymorphisms and cytosine
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                                                                                                                                                                                                                                                                                                                                            Claim 1; SEQ ID NO 193664; 29pp + Sequence Listing; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 9.4; DB 1; Length 13;
Pred. No. 1.3e+03;
0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 13 BP; 7 A; 2 C; 1 G; 3 T; 0 U; 0 Other;
                                                                    Berlin K;
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Matches 10; Conservative
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(EPIG-) EPIGENOMICS AG.
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Sequence 13 BP; 4 A; 5 C; 1 G; 3 T; 0 U; 0 Other;

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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nuclectide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF9989, ABH00010-ABF9989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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ftp.wipo.int/pub/published_pct_sequences
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Score 9.4; DB 1; Length 13;
Pred. No. 1.3e+03;
0; Mismatches 1; Indels
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Pred. No. 1.3e+03;
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Matches 10; Conservative
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Best Local Similarity
Matches 10; Conserv
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ABF99133 standard; DNA; 13 BP

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                                                                                                                                                                             SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; 88; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                               Oligonucleotide SEQ ID NO 199130 for detecting SNP TSC0049008.
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acid (DNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomers are used for diseases and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABE09989, ABF00010-ABE9989, ABH0010-ABE9989 and ABI00010-ABE82073 represent the oligomers described in the invention. NOTE: The sequence was obtained in electronic format from WIPO at the printed specification, but the wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                     This invention describes novel oligonucleotide primers or peptide nucleic
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                                                                                                                                                                                                                                                                                         German.
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                                                                                                                                                                                                                                                                                         Claim 1; SEQ ID NO 149155; 29pp + Sequence Listing;
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Best Local Similarity 90.9
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ABF49159/c
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nuclectide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardicvascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                           Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                               Claim 1; SEQ ID NO 149156; 29pp + Sequence Listing; German.
                                                                                                                                                                                                                                                                                                                                                                         12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03;
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WPI; 2001-657177/75
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central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABCO010-ABC9989, ABF00010-ABF9989, ABF00010-ABF9989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at fit, pub/published_pot_sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory,

Claim 1; SEQ ID NO 202888; 29pp + Sequence Listing; German.

methylation status.

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and oytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99899, ABF00010-ABF99989, ABH0010-ABF99989 and ABI00010-ABF82073 data for this patent did not form part of the printed specification, but was obtained in electronic formmat from WFPO at the printed specification, but the wipo.int/pub/published_pct_sequences
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                                                           SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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              Oligonuclectide SEQ ID NO 180118 for detecting SNP TSC0044592.
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Query Match

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ABF80121

ABF80121/ ID ABF8 XX AC ABF8 XX DT 22-F

RESULT 2095

Homo sapiens.

18-OCT-2001

olek A,

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Gaps

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schultz1-899.rng

SEQ ID NO 184331; 29pp + Sequence Listing; German.

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                                                                                                                                                                                                                                                                                                                                                        Claim 1; SEQ ID NO 205295; 29pp + Sequence Listing; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 12.9%; Score 9.4; DB 1; Length 13; Best Local Similarity 90.9%; Pred. No. 1.3e+03; Matches 10; Conservative 0; Mismatches 1; Indels
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06-APR-2001; 2001WO-IB000713.
                                                07-APR-2000; 2000DE-01019173.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABE99989, ABF00010-ABE99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.

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(first entry)

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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                      Oligonucleotide SEQ ID NO 213457 for detecting SNP TSC0051980.
                                                                      ABH13480 standard; DNA; 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                       12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03; ive 0; Mismatches 1; Indels
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                                                                         Sequence 13 BP; 2 A; 7 C; 0 G; 4 T; 0 U; 0 Other;
was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                                             10; Conservative
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Best Local Similarity 90.9
Matches 10, Conservative
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designed to detect single-nucleotide polymorphisms and cytosine
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ftp.wipo.int/pub/published_pct_sequences
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Local Similarity 90.9%;
Hes 10; Conservative C
                                                                                                      06-APR-2001; 2001WO-IB000713.
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                                                                                                                                                                                                                                                                 Piepenbrock C,
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                                                                                                                                                                                                                                                                                                                                                                                                                                    methylation status.
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ABH48201 standard;
WO200177384-A2.
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                                                18-OCT-2001.
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GTTTATTGTAT 11

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schultz1-899

Berlin K;

Olek A, Piepenbrock C,

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, contral nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABE99899, ABH00010-ABH99989 and ABI00010-ABI82073 tepresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at
central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                               Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                     Homo sapiens
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                                                                                                                                                                                                            Oligonucleotide SEQ ID NO 249230 for detecting SNP TSC0060878.
Query Match
12.9%; Score 9.4; DB 1; Length 13;
Best Local Similarity 90.9%; Pred. No. 1.3e+03;
Matches 10; Conservative 0; Mismatches 1; Indels
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                                                   947 GITTAATGTAT 957
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07-APR-2000; 2000DE-01019173. 06-APR-2001; 2001WO-IB000713

WO200177384-A2. Homo sapiens.

18-OCT-2001

(EPIG-) EPIGENOMICS AG

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but two botained in electronic format from WIPO at
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                 Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 12.9%; Score 9.4; DB 1; Length 13; Best Local Similarity 90.9%; Pred. No. 1.3e+03; Matches 10; Conservative 0; Mismatches 1; Indels
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                                                                                                                                                            Claim 1; SEQ ID NO 249230; 29pp + Sequence Listing; German
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Length 13;

Score 9.4; DB 1; Pred. No. 1.3e+03;

12.9%; 76.9%;

Query Match

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Matches
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and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF99899, ABF00010-ABF99989 and ABI0010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; 8s; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                       12.9%; Score 9.4; DB 1; Length 13; 76.9%; Pred. No. 1.3e+03; tive 1; Mismatches 2; Indels
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                                                                                                                                                                                                                    Sequence 13 BP; 1 A; 0 C; 3 G; 8 T; 0 U; 1 Other;
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Matches 10; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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designed to detect single-nucleotide polymorphisms and cytosine
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Pred. No. 1.3e+03;
0; Mismatches 1; Indels
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      1; Mismatches
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretraeted genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABH99989 and ABI00010-ABI82073 tapesent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but they wise obtained in electronic format from WIPO at
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                                                                                               SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                Oligonucleotide SEQ ID NO 96231 for detecting SNP TSC0023919.
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                                21-FEB-2002 (first entry)
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Best Local Similarity 90.9
Matches 10; Conservative
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nuclectide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, contral nervous system, axidovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 tepsesm the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at
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-ABC99989, ABF00010-ABF99989, ABH00010-ABH99889 and ABI00010-ABI82073 represent the oligomers described in the invention. NoTE: The Sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences

Sequence 13 BP; 2 A; 0 C; 3 G; 8 T; 0 U; 0 Other;

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastroinfrestinal, respitatory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010
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Matches 10; Conservative
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Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                           Claim 1; SEQ ID NO 75212; 29pp + Sequence Listing; German.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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Score 9.4; DB 1; Length 13;
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Matches 10; Conservative
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989 and ABI00010-ABF32073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo.int/pub/published_pct_sequences
                  SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                                                                                                                                 Claim 1; SEQ ID NO 103495; 29pp + Sequence Listing; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  March 12.9%; Score 9.4; DB 1; Length 13; Local Similarity 90.9%; Pred. No. 1.3e+03; es 10; Conservative 0; Mismatches 1; Indels
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07-APR-2000; 2000DE-01019173
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                                                                                                                                      Piepenbrock C,
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Claim 1; SEQ ID NO 103831; 29pp + Sequence Listing; German.

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligoners for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disconders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99999, ABC0010-ABC99999, ABC0010-ABC99999, ABC0010-ABC99999, ABC0010-ABC99999, ABC0010-ABC99999, ABC0010-ABC999999 and ABI00010-ABC99999 represent the oligomers described in the invention. NOTE: The sequence was obtained in electronic formm part of the printed specification, but the wipo.int/pub/published_pct_sequences
        This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010.ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a renge of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABR00010-ABF9989, ABH00010-ABH99899 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic form at from NIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                           SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; 88; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                                                                Oligonucleotide SEQ ID NO 111069 for detecting SNP TSC0027729.
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                       ABF11072 standard; DNA; 13 BP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                   Length 13;
                                                               Score 9.4; DB 1; Length 13
Pred. No. 1.3e+03;
0; Mismatches 1; Indels
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ABF09127 standard; DNA; 13 BP.
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         WPI; 2001-657177/75
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                                                                                                                                                                                                                                         This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99889, ABF00010-ABF99889 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                                                                                                                                            Score 9.4; DB 1; Length 13; Pred. No. 1.3e+03;
                                                                                                                                                                                                                      Claim 1; SEQ ID NO 16205; 29pp + Sequence Listing; German.
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                                                                                                                                Berlin K;
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Matches 10; Conservative
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                                                                                              Claim 1; SEQ ID NO 65343; 29pp + Sequence Listing; German
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range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                                                                                                                                                                                                                            Sequence 13 BP; 6 A; 4 C; 1 G; 2 T; 0 U; 0 Other;
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12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03;

Query Match Best Local Similarity

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Best Local Similarity
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Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine

methylation status.

Berlin K;

Piepenbrock C,

olek A,

WPI; 2001-657177/75

06-APR-2001; 2001WO-IB000713.

07-APR-2000; 2000DE-01019173

(EPIG-) EPIGENOMICS

Claim 1; SEQ ID NO 127230; 29pp + Sequence Listing; German.

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                                                                           SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                     Oligonucleotide SEQ ID NO 125011 for detecting SNP TSC0031240.
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21-FEB-2002 (first entry)
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF9989, ABF00010-ABF9989, ABF00010-ABF99989, ABF00010-ABF99989 and ABI00010-ABF82073 tapesees this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                    Claim 1; SEQ ID NO 134392; 29pp + Sequence Listing; German.
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Best Local Similarity 90.9
Matches 10; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                              SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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Pred. No. 1.3e+03;
0; Mismatches 1; Indels
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                                                                     Sequence 13 BP; 7 A; 3 C; 0 G; 3 T; 0 U; 0 Other;
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Best Local Similarity 90.9%;
Matches 10; Conservative
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les 10; Conservative
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07-APR-2000; 2000DE-01019173.
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                                                                                                                                                                                                                                                                                                                                                                                                                  This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC09989, ABF00010-ABF9989, ABH00010-ABH99899 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but typ.wipo.int/pub/published_pct_sequences
                                                                                                                               SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                           Oligonucleotide SEQ ID NO 197137 for detecting SNP TSC0048522.
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                                            ABF97140 standard; DNA; 13 BP.
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peptide nucleic acid, cytosine methylation, cardiovascular; primer; 88; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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designed to detect single-nucleotide polymorphisms and cytosine
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nuclectide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABE99899, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                          ligonucleotides, useful for diagnosis and cell typing, it o detect single-nucleotide polymorphisms and cytosine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Oligonucleotide SEQ ID NO 201920 for detecting SNP TSC0049639.
                                                                                                                                                   Claim 1; SEQ ID NO 173550; 29pp + Sequence Listing; German.
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                                                                                          oligonucleotides,
                             Olek A, Piepenbrock C,
EPIGENOMICS AG.
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acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically precreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory. Central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99889, ABC0010-ABC99899, ABC0010-ABC99899, ABC0010-ABC9989, ABC0010-ABC9980 and ABI00010-ABC8073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Pred. No. 1.3e+03;
0; Mismatches 1; Indels
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90.9%;
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This invention describes novel oligonuclectide primers or peptide nucleic

Claim 1; SEQ ID NO 201920; 29pp + Sequence Listing; German.

Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.

Berlin K;

Olek A, Piepenbrock C,

WPI; 2001-657177/75.

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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                  Oligonucleotide SEQ ID NO 228087 for detecting SNP TSC0055622.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 trepresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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            Score 9.4; DB 1; Length 13; Pred. No. 1.3e+03; 0; Mismatches 1; Indels
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                                                  This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and oytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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Claim 1; SEQ ID NO 228087; 29pp + Sequence Listing; German.
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Homo sapiens

ABH28110 standard; DNA; 13 BP.

RESULT 2134 ABH28110 ID ABH2811

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10; Conservative 908 TTTTCTTTGGT 918

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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretraeted genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, cancer also used for addiovacular and metabolic disorders. The oligomers are also used for adetecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF99899 and ABI00010-ABI82073 data for this patent did not form part of the printed specification, but two obtained in electronic format from WIPO at
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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 WO200177384-A2
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Berlin K;

WPI; 2001-657177/75.

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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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     designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                       Claim 1; SEQ ID NO 179260; 29pp + Sequence Listing; German.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, contral nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Best Local Similarity 90.9%; Pred. No. 1.3e+03;
Matches 10; Conservative 0; Mismatches 1; Indels
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Pred. No. 1.3e+03;
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Best Local Similarity 90.9%;
Matches 10; Conservative 0
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                       SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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designed to detect single-nucleotide polymorphisms and cytosine
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907 ATTITITIES
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07-APR-2000; 2000DE-01019173

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99899, ABH00010-ABH99989 and ABI00010-ABH32073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but they wipo int/pub/published_pct_sequences
                                    SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Oligonucleotide SEQ ID NO 158978 for detecting SNP TSC0040030.
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ABH41303;
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12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03; tive 0; Mismatches 1; Indels
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  Query Match
Best Local Similarity 90.9
Matches 10; Conservative
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RESULT 2141

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06-APR-2001; 2001WO-IB000713.

WO200177384-A2

18-OCT-2001

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABE99989, ABF00010-ABE99989, ABH00010-ABE99989 and ABI00010-ABI32073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo int/pub/published_pct_sequences
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                                              (EPIG-) EPIGENOMICS AG.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99999, ABF00010-ABF99999, ABH00010-ABH99999 and ABI00010-ABI82073 trepresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from MIPO at
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Claim 1; SEQ ID NO 241280; 29pp + Sequence Listing; German.
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Pred. No. 1.3e+03;
0; Mismatches 1; Indels
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nes 10; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                          SND; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                Score 9.4; DB 1; Length 13; Pred. No. 1.3e+03; 0; Mismatches 1; Indels
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                                Sequence 13 BP; 9 A; 0 C; 3 G; 1 T; 0 U; 0 Other;
ftp.wipo.int/pub/published_pct_sequences
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                                                                Query Match
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13 GGTTTTATGTA 3

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, contral nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99889, ABH00010-ABH99988 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at

Claim 1; SEQ ID NO 242135; 29pp + Sequence Listing; German.

Bonnror-899.rng

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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                  Oligonucleotide SEQ ID NO 253419 for detecting SNP TSC0061816.
             BP.
            ABH53442 standard; DNA; 13
                                     (first entry)
                                                                                                    WO200177384-A2
                                                                                         Homo sapiens
                                     22-FEB-2002
                         ABH53442;
RESULT 2145
      ABH53442
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18-OCT-2001.

07-APR-2000; 2000DE-01019173.

06-APR-2001; 2001WO-IB000713,

(EPIG-) EPIGENOMICS AG

Berlin K; Olek A, Piepenbrock C,

WPI; 2001-657177/75.

Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.

Claim 1; SEQ ID NO 253419; 29pp + Sequence Listing; German.

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF0010-ABE99989, ABF0010-ABE99989 and ABI0010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at

Sequence 13 BP; 2 A; 1 C; 2 G; 7 T; 0 U; 1 Other;

Gaps ö 12.9%; Score 9.4; DB 1; Length 13; 76.9%; Pred. No. 1.3e+03; ive 1; Mismatches 2; Indels Query Match
Best Local Similarity 76.9 Matches 10, Conservative

907 ATTITITITIGGIC 919

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ABH57691 standard; DNA; 13 BP ABH57691 2146 RESULT 21 ABH57691/ **EXEXEXEXEX**

Oligonucleotide SEQ ID NO 257668 for detecting SNP TSC0062683.

(first entry)

22-FEB-2002

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Homo sapiens.

WO200177384-A2

18-OCT-2001

06-APR-2001; 2001WO-IB000713

07-APR-2000; 2000DE-01019173

(EPIG-) EPIGENOMICS

Berlin K; Olek A, Piepenbrock C,

WPI; 2001-657177/75.

Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.

Claim 1; SEQ ID NO 257668; 29pp + Sequence Listing; German.

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABE99989, ABF00010-ABE99989, ABF00010-ABE99989, ABF00010-ABE99989, ABF00010-ABE99989 and ABI00010-ABE90010 data for this patent did not form part of the printed specification, but was obtained in electronic format from MIPO at

Sequence 13 BP; 8 A; 3 C; 0 G; 2 T; 0 U; 0 Other;

. 0 12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03; ive 0; Mismatches 1; Indels Query Match
Best Local Similarity 90.9

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908 TTTTCTTTGGT 918 m ||||| |||||| TTTTATTTGGT 13

ò a 2147 RESULT ABC93473 standard; DNA; 13

BP.

ABC93473;

21-FEB-2002 (first entry)

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Oligonucleotide SEQ ID NO 93490 for detecting SNP TSC0023360.

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Homo sapiens.

WO200177384-A2.

18-OCT-2001

06-APR-2001; 2001WO-IB000713

07-APR-2000; 2000DE-01019173.

(EPIG-) EPIGENOMICS

12.0

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Gaps

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Set of oligonucleotides, useful for diagnosis and cell typing, addesigned to detect single-nucleotide polymorphisms and cytosine
                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1; SEQ ID NO 95549; 29pp + Sequence Listing; German.
                                                                                                                                                                                                                                    ABC95532 standard; DNA; 13 BP.
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Best Local Similarity 76.9
Matches 10, Conservative
                         of oligonucleotides,
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Piepenbrock
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                                      methylation status
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretraeted genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, aradiovascular and metabolic diseoders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invantion. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at
                   range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The coligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99899, ABF00010-ABF99899, ABF00010-ABF99899 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
oligonucleotides are used for diagnosis and/or prognosis of cancer and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Oligonucleotide SEQ ID NO 97202 for detecting SNP TSC0024109
                                                                                                                                                                                                                                                                                                          12.9%; Score 9.4; DB 1; Length 13; 76.9%; Pred. No. 1.3e+03; ive 1; Mismatches 2; Indels
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                                                                                                                                                                                                                                                           Sequence 13 BP; 1 A; 1 C; 2 G; 8 T; 0 U; 1 Other;
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ABC97185/c
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                                                                                                                                                                                                                                                        This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nuclectide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form par of the printed specification, but was obtained in electronic format from WIPO at fire wipo.int/pub/published_pct_sequences
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Pred. No. 1.3e+03;
1; Mismatches 2; Indels
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typing, i cytosine

12.9%; Score 9.4; DB 1; Length 13;

Query Match

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The

Sequence 13 BP; 8 A; 2 C; 1 G; 2 T; 0 U; 0 Other;

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99999, ABF00010-ABF99989, ABH00010-ABH99999 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                   SNP; single nuclectide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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   Pred. No. 1.3e+03;
); Mismatches 1; Indels
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90.9%;
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Best Local Similarity 90.9
Matches 10; Conservative
 Best Local Similarity 90.9
Matches 10; Conservative
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; 85; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Oligonucleotide SEQ ID NO 2853 for detecting SNP TSC0001123.

20-FEB-2002

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Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.

Berlin K;

Piepenbrock C,

olek A,

WPI; 2001-657177/75

(EPIG-) EPIGENOMICS AG.

06-APR-2001; 2001WO-IB000713 07-APR-2000; 2000DE-01019173

WO200177384-A2.

18-OCT-2001

Homo sapiens.

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ABC27564 standard; DNA; 13
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Matches 10; Conservative
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This invention describes novel oligonucleotide primars or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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designed to detect single-nucleotide polymorphisms and cytosine
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                       methylation status.
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Pred. No. 1.3e+03;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                               Claim 1; SEQ ID NO 27581; 29pp + Sequence Listing; German
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                                               06-APR-2001; 2001WO-IB000713
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Best Local Similarity 90.9
Matches 10; Conservative
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                                                                                                                                                (EPIG-) EPIGENOMICS AG
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                                                                                                                                                                                                                                                                                                                                                methylation status.
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ABC28611

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Olek A,

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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                 ABC54364 standard; DNA; 13
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  3 TTTGGTTTTTG 13
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represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at the print pub/published_pct_sequences
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12.9%; Score 9.4; DB 1; Length 13;
Best Local Similarity 90.9%; Pred. No. 1.3e+03;
Matches 10; Conservative 0; Mismatches 1; Indels
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                                                                                                                            Sequence 13 BP; 6 A; 0 C; 1 G; 6 T; 0 U; 0 Other;
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ABC79370
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Berlin K;

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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 targresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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detect single-nucleotide polymorphisms and cytosine
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                                                                                                                                                                                                                                                                       Claim 1; SEQ ID NO 54928; 29pp + Sequence Listing; German.
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es 10; Conservative
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ATTTTTTTGTY 1
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABF3073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic formmat from WIPO at
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Local Similarity 90.9%; Pred. No. 1.38+03;
les 10; Conservative 0; Mismatches 1; Indels
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                                                                  Berlin K;
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(EPIG-) EPIGENOMICS AG.
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Sequence 13 BP; 6 A; 0 C; 5 G; 2 T; 0 U; 0 Other;

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99899, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but they wispo int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                                             Score 9.4; DB 1; Length 13; Pred. No. 1.3e+03; 0; Mismatches 1; Indels
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90.9%;
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Matches 10; Conservative
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                                                                                                                                                                                                                                                                                                                                SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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 12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03; Live 0; Mismatches 1; Indels
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Matches 10, Conservative
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Query Match
Best Local Similarity
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ABC62351 RESULT

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC09989, ABF00010-ABF9989, ABH00010-ABH9989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                           SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                Oligonucleotide SEQ ID NO 62368 for detecting SNP TSC0016537.
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ABC62351 standard; DNA; 13
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methylation status.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99889, ABF00010-ABF99899, ABH00010-ABF99899 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic formmat from NIPO at fitte.wipo.int/pub/published_pct_sequences
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Matches 10; Conservative
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretraeted genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF9989, ABH00010-ABF9989 and ABI00010-ABF8073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                  Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine
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                                                                                                                                                                                                                                                                                                                                                                                                                                              12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03; ive 0; Mismatches 1; Indels
                                                                                                               Claim 1; SEQ ID NO 90623; 29pp + Sequence Listing; German.
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Best Local Similarity 90.9
Matches 10, Conservative
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WPI; 2001-657177/75
                                                                            methylation status.
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central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99889, ABF00010-ABF9989, ABF00010-ABF9989, ABF00010-ABF9989, ABF00010-ABF9989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this parent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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90.9%;
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Best Local Similarity 90.9%;
Matches 10; Conservative
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Oligonucleotide SEQ ID NO 125012 for detecting SNP TSC0031240.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; 8s; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Pred. No. 1.3e+03;
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, acadiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99999, ABF00010-ABF99989, ABF00010-ABF9989, ABF00010-ABF89989, ABF00010-ABF89989 and ABF00010-ABF8073 are present the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from MIPO at
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                                           SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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06-APR-2001; 2001WO-IB000713.
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RATCTATCCCTAC 1
                                                                                      Olek A, Piepenbrock C,
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                                                         (EPIG-) EPIGENOMICS AG
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                                                                                                                    WPI; 2001-657177/75.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, cantral nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99999, ABF00010-ABF99999, ABH00010-ABH99999 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This invention describes novel oligonucleotide primers or peptide nucleic and (PMA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, oligomers are also used for detecting cell type differentiation. ABC00010 eligomers are also used for detecting cell type differentiation. ABC0001 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Claim 1; SEQ ID NO 136951; 29pp + Sequence Listing; German.
                                                                                                                                                                                                                                                                                                                                                                                                    12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 1; SEQ ID NO 139201; 29pp + Sequence Listing; German.
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                                                                                     Oligonucleotide SEQ ID NO 218935 for detecting SNP TSC0053255.
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                                                      ABH18958 standard; DNA; 13
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                  RESULT 2173
ABH18958/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                          Score 9.4; DB 1; Length 13; Pred. No. 1.3e+03; 0; Mismatches 1; Indels
                                                        Sequence 13 BP; 5 A; 0 C; 3 G; 5 T; 0 U; 0 Other;
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was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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                                                                                                  12.9%;
90.9%;
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Matches 10, Conservative
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                                                                                                                                          Matches
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ABP43154

ABP43154

ABP7

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically prereated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABC0010-ABE9989, ABH00010-ABE99989, ABH00010-ABE99989, ABH00010-ABE99989, and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but twipo.int/pub/published_pot_sequences
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Pred. No. 1.3e+03;
0; Mismatches 1; Indels
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3 ATTTTTTTGG 13

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Berlin K;

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Piepenbrock C,
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            olek A,
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ABH23858
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                                                                                                                                                                                                                                                                                                             This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
central nervous system; gastrointestinal; respiratory; immune; metabolic
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Pred. No. 1.3e+03;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 13 BP; 1 A; 0 C; 2 G; 10 T; 0 U; 0 Other;
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                                                                                                                                                                                 Berlin K;
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Best Local Similarity 90.9
Matches 10; Conservative
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                                                WO200177384-A2
                          Homo sapiens.
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(EPIG-) EPIGENOMICS AG

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABE99989, ABF00010-ABE99989, ABH00010-ABE99989 and ABI00010-ABE82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                         Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine
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                                                                                                                                     Claim 1; SEQ ID NO 171263; 29pp + Sequence Listing; German.
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WPI; 2001-657177/75
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and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABC9989, ABH00010-ABC9989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nuclectide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form par to f the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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designed to detect single-nucleotide polymorphisms and cytosine
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Seguence 13 BP; 1 A; 1 C; 3 G; 8 T; 0 U; 0 Other;

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Score 9.4; DB 1; Length 13; Pred. No. 1.3e+03;
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF99989, ABH00010-ABF99989, ABH00010-ABF99989, ABH0010-ABF99989 and ABI00010-ABF8003 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo.int/pub/published_pot_sequences
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                                                                                                                                                                                          SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                            Oligonucleotide SEQ ID NO 201919 for detecting SNP TSC0049639.
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                                                        methylation status.
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ABP754515
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic formmat from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at the published_pot_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                                        Sequence 13 BP; 4 A; 4 C; 1 G; 4 T; 0 U; 0 Other;
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Matches 10, Conservative
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                                                                                                                                                     This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF99989, ABH00010-ABF99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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  oligonucleotides, useful for diagnosis and cell typing, :
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                                                                                                               Claim 1; SEQ ID NO 179259; 29pp + Sequence Listing; German.
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Pred. No. 1.3e+03;
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                                                                                                                    RESULT 2184
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ABF84447/
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nuclectide polymorphisms (SNP) and cytosine methylation status in chemically pretraeted genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic discorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABE99989, ABF00010-ABE99989, ABH00010-ABE99989 and ABI00010-ABI32073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WFPO at
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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07-APR-2000; 2000DE-01019173

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABE9989, ABF00010-ABE9989, and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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Claim 1; SEQ ID NO 236754; 29pp + Sequence Listing; German

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    This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, contral nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from MIPO at
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designed to detect single-nuclectide polymorphisms and cytosine
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Oligonucleotide SEQ ID NO 241279 for detecting SNP TSC0058852.

(first entry)

22-FEB-2002

ABH41302;

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ABH41302 standard; DNA; 13

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                                                                                                                                                                                                                                                                          SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                     Length 13;
                                  Score 9.4; DB 1; Length 13
Pred. No. 1.3e+03;
0; Mismatches 1; Indels
            Sequence 13 BP; 4 A; 0 C; 2 G; 7 T; 0 U; 0 Other;
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Matches 10; Conservative
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at

Sequence 13 BP; 5 A; 0 C; 4 G; 4 T; 0 U; 0 Other;

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set or oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.

Berlin K;

Olek A, Piepenbrock C,

WPI; 2001-657177/75.

(EPIG-) EPIGENOMICS AG.

07-APR-2000; 2000DE-01019173. 06-APR-2001; 2001WO-IB000713

40200177384-A2. Homo sapiens

18-OCT-2001

Claim 1; SEQ ID NO 241279; 29pp + Sequence Listing; German.

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Query Match
12.9%; Score 9.4; DB 1; Length 13;
Best Local Similarity 90.9%; Pred. No. 1.3e+03;
Matches 10; Conservative 0; Mismatches 1; Indels
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF9989, ABH0010-ABH99899 and AB100010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from MIPO at
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                              ber or oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                           Claim 1; SEQ ID NO 251783; 29pp + Sequence Listing; German.
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Pred. No. 1.3e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                                                                                                                      Claim 1; SEQ ID NO 245088; 29pp + Sequence Listing; German.
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    Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretraeted genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form par of the printed specification, but ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Best Local Similarity 90.9
Matches 10; Conservative
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                                                                                                                                                                                                                                                      SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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12.9%; Score 9.4; DB 1; Length 13;
Best Local Similarity 90.9%; Pred. No. 1.3e+03;
Matches 10; Conservative 0; Mismatches 1; Indels
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AC ABC70879;
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                       TTTATCCTTCC 12
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927 ITTAICCTCC
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Length 13;

Score 9.4; DB 1; Pred. No. 1.3e+03;

12.9%;

Query Match Best Local Similarity

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                                                                          SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                         Oligonucleotide SEQ ID NO 70896 for detecting SNP TSC0018403.
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Matches 10; Conser
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ABP00359
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AC ABF00355
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WO200177384-A2

18-OCT-2001.

Homo sapiens

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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99889, ABF00010-ABF99989, ABH00010-ABF99989, ABH0010-ABF99989 and ABI00010-ABF9073 represent the oligomers described in the invention. NOTE: The sequence was obtained in electronic format from WIPO at the printed specification, but the wipo.int/pub/published_pct_sequences
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90.9%;
06-APR-2001; 2001WO-IB000713.
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                                                                                                                                 This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, contral nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                    Claim 1; SEQ ID NO 76038; 29pp + Sequence Listing; German.
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                                                                                                                    Seguence 13 BP; 1 A; 8 C; 0 G; 4 T; 0 U; 0 Other
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was obtained in electronic format from WIPO at
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peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

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Piepenbrock C,

olek A,

(EPIG-) EPIGENOMICS AG

06-APR-2001; 2001WO-IB000713. 07-APR-2000; 2000DE-01019173.

WO200177384-A2.

18-OCT-2001

Homo sapiens.

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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS;
                                                                                                                                       Oligonucleotide SEQ ID NO 103950 for detecting SNP TSC0025999
                                                                         ABF03953 standard; DNA; 13 BP.
945 TGGTTTAATGTAT 957
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99889, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but fyp. wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Pred. No. 1.3e+03;
1; Mismatches 2; Indels
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                                                                                         ABF02167 standard; DNA; 13
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Best Local Similarity 76.99
Marches 10; Conservative
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RESULT 2201
ABF02167/C
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABF99989 and ABI00010-ABF32073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo.int/pub/published_pot_sequences
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                                                                                                                                                                                                                                                                                                                                                 Claim 1; SEQ ID NO 103950; 29pp + Sequence Listing; German.
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76.9%;
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Best Local Similarity 76.99
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                                                                                                                                                                                                       This invention describes novel oligonucleotide primers or peptide nucleic
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                                Berlin K;
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                                Piepenbrock C,
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 (EPIG-) EPIGENOMICS AG.
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Best Local Similarity
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acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically precreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastronintestinal, respiratory. Central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99889, ABC0010-ABC99889, ABC0010-ABC99889, ABC0010-ABC9988, and ABI00010-ABC8073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF9989, ABH00010-ABF9989 and ABI00010-ABF9073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Pred. No. 1.3e+03;
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Best Local Similarity 90.9%;
Matches 10; Conservative
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

WO200177384-A2.

Oligonucleotide SEQ ID NO 37840 for detecting SNP TSC0011749.

(first entry)

20-FEB-2002

ABC37823;

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                                                                                                                                                                                                                                                                                                                                SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                      Gaps
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Score 9.4; DB 1; Length 13; Pred. No. 1.3e+03; 1; Mismatches 2; Indels
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ABC37822 standard; DNA; 13 BP.
 12.9%;
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Best Local Similarity 76.9
Matches 10; Conservative
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Best Local Similarity 90.9
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Pred. No. 1.3e+03;
0; Mismatches 1; Indels
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90.9%;
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ABC37823 standard; DNA; 13 BP.

RESULT 2207 ABC37823 ID ABC37823

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                    Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 9.4; DB 1; Length 13;
Pred. No. 1.3e+03;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                 Claim 1; SEQ ID NO 88590; 29pp + Sequence Listing; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 13 BP; 7 A; 2 C; 0 G; 3 T; 0 U; 1 Other;
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                                                                                                                                                          Berlin K;
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90.9%;
                                                             06-APR-2001; 2001WO-IB000713.
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Best Local Similarity
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WO200177384-A2
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acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99899, ABC0010-ABC99899 ABH0010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at the printed specification, but fire wipo.int/pub/published_pct_sequences
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                                                                                                                                     This invention describes novel oligonucleotide primers or peptide nucleic
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            Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine methylation status:
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                                                                                             Claim 1; SEQ ID NO 113693; 29pp + Sequence Listing; German.
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ABC39013 standard; DNA; 13 BP.
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oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF9989, ABF00010-ABF9989, ABF00010-ABF9989, and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                       SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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designed to detect single-nucleotide polymorphisms and cytosine
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Best Local Similarity 90.9%; Pred. No. 1.3e+03;
Matches 10; Conservative 0; Mismatches 1; Indels
                                                                                                                             Match 12.9%; Score 9.4; DB 1; Length 13; Local Similarity 90.9%; Pred. No. 1.3e+03; les 10; Conservative 0; Mismatches 1; Indels
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                                                                                                    Sequence 13 BP; 9 A; 3 C; 0 G; 1 T; 0 U; 0 Other;
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic formmat from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                      SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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designed to detect single-nucleotide polymorphisms and cytosine
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                                                                                                                                                                                                                                                                                                                                               Oligonucleotide SEQ ID NO 65741 for detecting SNP TSC0017295.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABF3073 data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                       oet or oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                                                                                                                                                                                                               Claim 1; SEQ ID NO 120915; 29pp + Sequence Listing; German.
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                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Oligonucleotide SEQ ID NO 119568 for detecting SNP TSC0029845.
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ABF20918 standard; DNA; 13
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genemic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH999989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                        This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABH999989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from MIPO at
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Claim 1; SEQ ID NO 126354; 29pp + Sequence Listing; German.
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Sequence 13 BP; 7 A; 4 C; 1 G; 0 T; 0 U; 1 Other;
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Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                   Oligonucleotide SEQ ID NO 143100 for detecting SNP TSC0035891.
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                        ABF43103 standard; DNA; 13
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABF99989, ABH00010-ABH99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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Berlin K;

Piepenbrock C,

olek A,

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The
                                                                                                                                                                   This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nuclectide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99999, ABF00010-ABF99999, ABH00010-ABH99999 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                          Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                      Claim 1; SEQ ID NO 143682; 29pp + Sequence Listing; German
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                                WPI; 2001-657177/75
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Best Local Similarity
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancr and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99389, ABC0010-ABC99389, ABC0010-ABC99389, ABC0010-ABC99389, ABC0010-ABC99389, ABC0010-ABH99999 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence was obtained in electronic form part of the printed specification, but frp.wipo.int/pub/published_pct_sequences
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              range of diseases including immune system, gastrointestinal, respiratory, range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting call type differentiation. ABC00010-ABC099889, ABF00010-ABH99889 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at fire wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SNP, single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
used for diagnosis and/or prognosis of cancer and
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Pred. No. 1.3e+03;
0; Mismatches 1; Indels
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Best Local Similarity 90.9
Matches 10, Conservative
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12.9%; Score 9.4; DB 1; Length 13;

Query Match

Sequence 13 BP; 1 A; 8 C; 1 G; 3 T; 0 U; 0 Other;

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABE99989, ABF00010-ABE99989, ABH00010-ABE99989 and ABI00010-ABE82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                 SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                       Oligonucleotide SEQ ID NO 173291 for detecting SNP TSC0043175
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Best Local Similarity
Matches 10; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                   SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic
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12.9%; Score 9.4; DB 1; Length 13;
Best Local Similarity 90.9%; Pred. No. 1.3e+03;
Matches 10; Conservative 0; Mismatches 1; Indels
                        Indels
  Pred. No. 1.3e+03;
0; Mismatches 1;
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ftp.wipo.int/pub/published_pct_sequences
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Best Local Similarity 90.9%;
Matches 10; Conservative
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ID ABF7
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AC ABF7

06-APR-2001; 2001WO-IB000713.

18-OCT-2001.

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99999, ABF00010-ABF99999, ABH00010-ABH99999 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99999, ABF00010-ABF99999, ABH00010-ABH99999 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic formmat from WIPO at
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designed to detect single-nucleotide polymorphisms and cytosine
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                                                        claim 1; SEQ ID NO 149516; 29pp + Sequence Listing; German.
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Pred. No. 1.3e+03;
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Local Similarity 90.9%;
les 10; Conservative C
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                      methylation status.
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ABF78025 standard; DNA; 13
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represent the oligomers described in the invention. NOTE: The sequence data for this parent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences 12.9%; Score 9.4; DB 1; Length 13; llarity 76.9%; Pred. No. 1.38+03; Conservative 1; Mismatches 2; Indels Sequence 13 BP; 6 A; 2 C; 0 G; 4 T; 0 U; 1 Other; Query Match Best Local Similarity Matches 10; Conserv

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943 ATTGGTTTAATGT 955

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RESULT 2228

BP ABH00850 standard; DNA; 13

ABH00850;

22-FEB-2002 (first entry)

Oligonucleotide SEQ ID NO 200827 for detecting SNP TSC0049410.

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Homo sapiens.

WO200177384-A2.

18-OCT-2001

06-APR-2001; 2001WO-IB000713

07-APR-2000; 2000DE-01019173

(EPIG-) EPIGENOMICS AG.

Olek A, Piepenbrock C,

× Berlin

WPI; 2001-657177/75.

oligonuclectides, useful for diagnosis and cell typing, ied to detect single-nuclectide polymorphisms and cytosine methylation status. Set of

Claim 1; SEQ ID NO 200827; 29pp + Sequence Listing; German.

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo int/pub/published_pct_sequences

940 TICATIGGITT 950

Sequence 13 BP; 1 A; 0 C; 2 G; 9 T; 0 U; 1 Other;

Gaps ., Query Match 12.9%; Score 9.4; DB 1; Length 13; Best Local Similarity 90.9%; Pred. No. 1.38+03; Matches 10; Conservative 0; Mismatches 1; Indels

ABF78025;

(first entry) 22-FEB-2002

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Gaps

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Oligonucleotide SEQ ID NO 178022 for detecting SNP TSC0044112

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Homo sapiens

WO200177384-A2.

18-OCT-2001

06-APR-2001; 2001WO-IB000713.

07-APR-2000; 2000DE-01019173.

(EPIG-) EPIGENOMICS AG.

Berlin Olek A, Piepenbrock C,

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WPI; 2001-657177/75.

Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.

claim 1; SEQ ID NO 178022; 29pp + Sequence Listing; German

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at

Sequence 13 BP; 6 A; 1 C; 0 G; 5 T; 0 U; 1 Other;

Gaps ; Query Match 12.9%; Score 9.4; DB 1; Length 13; Best Local Similarity 76.9%; Pred. No. 1.3e+03; Matches 10; Conservative 1; Mismatches 2; Indels

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ઠ g RESULT 2230
ABF80153
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AC ABF80153
AC ABF80155
DT 22-FEB-2
XX

ABF80153 standard;

DNA; 13

ABF80153;

22-FEB-2002 (first entry)

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Oligonucleotide SEQ ID NO 180150 for detecting SNP TSC0044601.

19. 13 · 1 da Cor German.

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic formmat from WIPO at
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                                                                                                              Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine methylation status.
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designed to detect single-nucleotide polymorphisms and cytosine
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                                                                                                                                                                                Claim 1; SEQ ID NO 205519; 29pp + Sequence Listing;
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                                                Piepenbrock C,
                 (EPIG-) EPIGENOMICS AG.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SNP, single nucleotide polymorphism, human; diagnosis; PNA, cancer; CNS, peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03; rive 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                        designed to detect
methylation status.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99899, ABF00010-ABF99899 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo.int/pub/published_pct_sequences

Sequence 13 BP; 3 A; 3 C; 0 G; 6 T; 0 U; 1 Other;

Gaps .; 0 Length 13; 2; Indels Query Match
12.9%; Score 9.4; DB 1;
Best Local Similarity 76.9%; Pred. No. 1.3e+03;
Matches 10; Conservative 1; Mismatches 2; 923 GCCITTIATCCCI 935 셤 ò

ABF80831 standard; DNA; 13 RESULT 2233

ВР.

Oligonucleotide SEQ ID NO 180828 for detecting SNP TSC0044744. (first entry) 22-FEB-2002

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Homo sapiens

WO200177384-A2.

18-OCT-2001,

06-APR-2001; 2001WO-IB000713.

07-APR-2000; 2000DE-01019173.

(EPIG-) EPIGENOMICS AG

Olek A, Piepenbrock C, WPI; 2001-657177/75.

Berlin K;

Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.

Claim 1; SEQ ID NO 180828; 29pp + Sequence Listing; German.

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF99899 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence was obtained in electronic format from WIPO at the printed specification, but the wipo.int/pub/published_pct_sequences

Gaps ., Length 13; 1, Indels Sequence 13 BP; 6 A; 2 C; 0 G; 5 T; 0 U; 0 Other; Score 9.4; DB 1; Pred. No. 1.3e+03; 0; Mismatches 1; 12.9%; 90.9%; Conservative Query Match Best Local Similarity Matches 10; Conserv g

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ABF81513 standard; DNA; 13 BP

ABF81513;

22-FEB-2002 (first entry)

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Oligonucleotide SEQ ID NO 181510 for detecting SNP TSC0044883.

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Homo sapiens.

WO200177384-A2.

18-OCT-2001

06-APR-2001; 2001WO-IB000713

07-APR-2000; 2000DE-01019173

(EPIG-) EPIGENOMICS AG.

WPI; 2001-657177/75.

Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.

This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nuclectide polymorphisms (SNP) and cytosine methylation status in chemically prereated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic discorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99889, ABF00010-ABF99889, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence and not this patent did not form part of the printed specification, but the wipo.int/pub/published_pct_sequences 8XXCCCCCCCCCCX8X44X4X4X4X4X6X6X6X8X6X6XXXXXXX8X8

Sequence 13 BP; 8 A; 2 C; 0 G; 2 T; 0 U; 1 Other;

Query Match 12.9%; Score 9.4; DB 1; Length 13; Best Local Similarity 76.9%; Pred. No. 1.3e+03; Matches 10; Conservative 1; Mismatches 2; Indels

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Gaps

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TCTTCATTGGTTT 950 | || || || || || || || || TTTTAATTGGTTY 1 938 13

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RESULT 2235 ABH08284/c

DITT: 660 T91TNI

Berlin K; Olek A, Piepenbrock C,

Claim 1; SEQ ID NO 181510; 29pp + Sequence Listing; German.

Berlin K;

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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and oytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                             Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                                                                                                                                                                                                                                                                                                    Claim 1; SEQ ID NO 208896; 29pp + Sequence Listing; German
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12.9%; Score 9.4; DB 1; Length 13; 76.9%; Pred. No. 1.38+03; vative 1; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 13 BP; 9 A; 3 C; 0 G; 0 T; 0 U; 1 Other;
                                                                                                 06-APR-2001; 2001WO-IB000713.
                                                                                                                                          07-APR-2000; 2000DE-01019173.
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es 10; Conservative
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                                                                                                                                                                                 (EPIG-) EPIGENOMICS AG
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                   WO200177384-A2
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ABH10811/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABC0010-ABE99899, ABC0010-ABE99899 and ABI0010-ABE90013 represent the oligomers described in the invention. NOTE: The sequence and act for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at the printed specification, but ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                         SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; SS; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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designed to detect single-nucleotide polymorphisms and cytosine
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                                                                                                                       Oligonucleotide SEQ ID NO 208261 for detecting SNP TSC0050910.
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    ABH08284 standard; DNA; 13
                                                                                  (first entry)
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                                           ABH08284
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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ABH10811 standard; DNA; 13 BP.
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Page 979

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretraeted genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at
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                                        Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                              Claim 1; SEQ ID NO 210788; 29pp + Sequence Listing; German.
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Best Local Similarity 90.9
Matches 10, Conservative
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WPI; 2001-657177/75
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic. Oligonucleotide SEQ ID NO 211394 for detecting SNP TSC0051569. BP. 417/c ABH11417 standard; DNA; 13 (first entry) 22-FEB-2002 ABH11417;

ABH11417,

WO200177384-A2 18-OCT-2001.

Homo sapiens

06-APR-2001; 2001WO-IB000713. 07-APR-2000; 2000DE-01019173.

(EPIG-) EPIGENOMICS AG

Berlin K; Olek A, Piepenbrock C,

WPI; 2001-657177/75

Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.

Claim 1; SEQ ID NO 211394; 29pp + Sequence Listing; German.

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory,

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central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99999, ABF00010-ABE99989 and ABI00010-ABI392073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                    Sequence 13 BP; 8 A; 2 C; 0 G; 2 T; 0 U; 1 Other;
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Best Local Similarity 90.9%;
Matches 10; Conservative
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ABF62509 standard; DNA; 13
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Oligonucleotide SEQ ID NO 217021 for detecting SNP TSC0052748.
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                                                                                                                                                                                                                                                                            methylation status.
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ABH51807/
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                                                                                                                                                                          SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                      Oligonuclectide SEQ ID NO 239884 for detecting SNP TSC0008514.
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ABH17044
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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designed to detect single-nucleotide polymorphisms and cytosine
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Pred. No. 1.3e+03;
1; Mismatches 2; Indels
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Best Local Similarity 76.99
Matches 10, Conservative
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                                                                                                                                                                                                                                                                                                          Claim 1; SEQ ID NO 251784; 29pp + Sequence Listing; German.
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Pred. No. 1.3e+03;
0; Mismatches 1; Indels
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                                                                                                                                 Berlin K;
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06-APR-2001; 2001WO-IB000713
                                        07-APR-2000; 2000DE-01019173
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Best Local Similarity 90.9%;
Matches 10; Conservative
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting call type differentiation. ABC0010 ABC99989, ABF00010-ABF99899 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo.int/pub/published_pct_sequences
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Claim 1; SEQ ID NO 253874; 29pp + Sequence Listing; German.
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Best Local Similarity
Matches 10; Conserv
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF9989, ABH00010-ABH9989 and ABI00010-ABI82073 trepresent the oligomers described in the invention. NOTE: The sequence date for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                   SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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designed to detect single-nucleotide polymorphisms and cytosine
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ftp.wipo.int/pub/published_pct_sequences
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                                                              ABH56217 standard; DNA; 13 BP.
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Pred. No. 1.3e+03;
0; Mismatches 1; Indels
                                                                                                         12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03; ive 0; Mismatches 1; Indels
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                                                                Sequence 13 BP; 6 A; 3 C; 0 G; 4 T; 0 U; 0 Other;
was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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90.9%;
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Matches 10; Conservative
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                                                                                                                                     Local Similarity
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Query Match

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Piepenbrock C,

olek A,

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, contral nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99889 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at
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central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Matches 10; Conservative
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but twipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                         Claim 1; SEQ ID NO 94182; 29pp + Sequence Listing; German.
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Matches 10; Conservative
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Score 9.4; DB 1; Length 13 Pred. No. 1.3e+03; 0; Mismatches 1; Indels

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Query Match Best Local S: Matches 10

946 GGTTTAATGTA 956

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99899, ABF00010-ABE99899, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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designed to detect single-nucleotide polymorphisms and cytosine
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                                                                                                                                                              SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; Ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Pred. No. 1.3e+03;
0; Mismatches 1; Indels
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                                 ABC21177 standard; DNA; 13 BP.
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ABC71594 standard; DNA; 13 BP.

ABC71594 ID ABC7 XX

Sequence 13 BP; 3 A; 1 C; 3 G; 6 T; 0 U; 0 Other;

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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                                                                                                                                                                                oet or oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                                                                                                                                                                                                                                                                              Claim 1; SEQ ID NO 71631; 29pp + Sequence Listing; German.
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                                                                                                                                                                                                           Berlin K;
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                                                                       06-APR-2001; 2001WO-IB000713.
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                                                                                                                                    SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                          Oligonucleotide SEQ ID NO 71611 for detecting SNP TSC0018532.
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                                         21-FEB-2002 (first entry)
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Best Local Similarity 76.5.
Best Local 10, Conservative
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ABC71594;
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RESULT 2253

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represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences

-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073

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acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
of oligonucleotides, useful for diagnosis and cell typing, . igned to detect single-nucleotide polymorphisms and cytosine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Oligonucleotide SEQ ID NO 49360 for detecting SNP TSC0013972.
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Pred. No. 1.3e+03;
1; Mismatches 2; Indels
                                                        Claim 1; SEQ ID NO 23961; 29pp + Sequence Listing; German
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les 10; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                     SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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designed to detect single-nucleotide polymorphisms and cytosine
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                                                                                                                        Score 9.4; DB 1; Length 13; Pred. No. 1.3e+03; 0; Mismatches 1; Indels
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                                                                                            Sequence 13 BP; 7 A; 4 C; 0 G; 1 T; 0 U; 1 Other;
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Best Local Similarity 90.9
Matches 10, Conservative
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Matches

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for dagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010

of oligonucleotides, useful for diagnosis and cell typing, igned to detect single-nucleotide polymorphisms and cytosine

Claim 1; SEQ ID NO 49360; 29pp + Sequence Listing; German.

methylation status.

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WO200177384-A2.
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                                                                                                                                                                                                                                                                                                                                                   SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99999, ABF00010-ABF99999, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Olek A,

WO200177384-A2. Homo sapiens

ABC78032;

Query Match

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07-APR-2000; 2000DE-01019173

Berlin K; Piepenbrock C, WPI; 2001-657177/75. olek A,

of oligonucleotides, useful for diagnosis and cell typing, igned to detect single-nucleotide polymorphisms and cytosine methylation status. Set of oldesigned

Claim 1; SEQ ID NO 78049; 29pp + Sequence Listing; German.

ABC04590 standard; DNA; 13 BP. 06-APR-2001; 2001WO-IB000713. 07-APR-2000; 2000DE-01019173. 20-FEB-2002 (first entry) 918 2 Trrrarrregr 12 (EPIG-) EPIGENOMICS AG 908 TITICITIGGI WO200177384-A2. Homo sapiens. 18-OCT-2001. ABC04590; RESULT 2261 *55555555555555 ઠ 8 ő This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic. Gaps oligonucleotides, useful for diagnosis and cell typing, is to detect single-nucleotide polymorphisms and cytosine ö Oligonucleotide SEQ ID NO 78049 for detecting SNP TSC0019867. Claim 1; SEQ ID NO 102163; 29pp + Sequence Listing; German. 12.9%; Score 9.4; DB 1; Length 13; 76.9%; Pred. No. 1.3e+03; tive 1; Mismatches 2; Indels Sequence 13 BP; 3 A; 0 C; 5 G; 4 T; 0 U; 1 Other; ftp.wipo.int/pub/published_pct_sequences Berlin K; ABC78032 standard; DNA; 13 BP 07-APR-2000; 2000DE-01019173 945 TGGTTTAATGTAT 957 21-FEB-2002 (first entry) 1 TGGTGTAATGGAY 13 Local Similarity 76.5 Piepenbrock C, (EPIG-) EPIGENOMICS AG WPI; 2001-657177/75 methylation status.

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABF32073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at ö This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABE03989, ABE00010-ABE9989, ABE00010-ABE9989, ABE00010-ABE9989, and ABI00010-ABE32073 represent the oligomers described in the invention. NOTE: The sequence was obtained in electronic format from WIPO at the printed specification, but typ.wipo.int/pub/published_pct_sequences SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic. Gaps onuclectides, useful for diagnosis and cell typing, idetect single-nuclectide polymorphisms and cytosine ; 0 Oligonuclectide SEQ ID NO 4581 for detecting SNP TSC0001664. Query Match
Best Local Similarity 90.9%; Pred. No. 1.3e+03;
Matches 10; Conservative 0; Mismatches 1; Indels Claim 1; SEQ ID NO 4581; 29pp + Sequence Listing; German. Seguence 13 BP; 1 A; 0 C; 2 G; 10 T; 0 U; 0 Other; Set of oligonucleotides, designed to detect single Piepenbrock C, WPI; 2001-657177/75. methylation status. olek A,

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99899, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo.int/pub/published_pot_sequences
                                                                                                                                           SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                           Oligonucleotide SEQ ID NO 11202 for detecting SNP TSC0002751.
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   ABC11211 standard; DNA; 13 BP.
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Sequence 13 BP; 2 A; 0 C; 3 G; 8 T; 0 U; 0 Other;
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABF99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIFO at
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                                                                                oligonucleotides, useful for diagnosis and cell typing, ied to detect single-nucleotide polymorphisms and cytosine
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                                                                                                                                                                                                                                                                                                                                                                      Berlin K;
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range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC09989, ABF00010-ABF9988, ABF00010-ABF9989 ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form par of the printed specification, but was obtained in electronic format from WIPO at the printed specification, but flow wipo.int/pub/published_pct_sequences
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at fitte.wipo.int/pub/published_pot_sequences
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                                                                                                                                                                                                                  Oligonucleotide SEQ ID NO 122312 for detecting SNP TSC0030569.
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Length 13;

12.9%; Score 9.4; DB 1; 90.9%; Pred. No. 1.3e+03;

Query Match Best Local Similarity

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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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designed to detect single-nucleotide polymorphisms and cytosine
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                                                                                           SNP, single nucleotide polymorphism, human; diagnosis; PNA; cancer, CNS, peptide nucleic acid; cytosine methylation; cardiovascular; primer; 88; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                               Oligonucleotide SEQ ID NO 131636 for detecting SNP TSC0032855.
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Pred. No. 1.3e+03;
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Matches 10; Conservative
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
Oligonucleotide SEQ ID NO 139202 for detecting SNP TSC0034868
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                                                        This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form mar of the printed specification, but was obtained in electronic format from WIPO at the printed specification, but firm wipo.int/pub/published_pct_sequences
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                            Claim 1; SEQ ID NO 139202; 29pp + Sequence Listing; German.
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ABH18708 standard; DNA; 13
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Matches 10; Conservative
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methylation status
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data for this patent did not form part of the pass obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                        SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                     Oligonucleotide SEQ ID NO 195989 for detecting SNP TSC0048213.
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                                                                                       ABF95992 standard; DNA; 13 BP.
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peptide nucleic acid, cytosine methylation; cardiovascular; primer; 88; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                      Claim 1; SEQ ID NO 198715; 29pp + Sequence Listing; German.
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acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory. Central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99889, ABC0010-ABC9989, ABH0010-ABC9989, ABH0010-ABC9989, ABH0010-ABC9989 and ABI00010-ABC8073 data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

WO200177384-A2.

18-OCT-2001.

Homo sapiens

Oligonucleotide SEQ ID NO 177488 for detecting SNP TSC0044012.

(first entry)

22-FEB-2002

ABF77491;

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12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03; ive 0; Mismatches 1; Indels
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Pred. No. 1.3e+03;
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Homo sapiens

ABF77491 standard; DNA; 13

RESULT 2280 ABF77491/c ID ABF77491

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but ftp.wipo.int/pub/published_pct_sequences
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Matches 10; Conservative
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting call type differentiation. ABC00010-ABC9989, ABF00010-ABF9989, ABH00010-ABH9989 and ABI00010-ABI32073 carepresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                  Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                                                              Claim 1; SEQ ID NO 203270; 29pp + Sequence Listing; German.
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ftp.wipo.int/pub/published_pct_sequences
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oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF9989, ABH00010-ABH9989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at fibe.wipo.int/pub/published_pct_sequences
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABF99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                                                RESULT 2285
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ID ABF8
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AC ABF8
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22-FEB-2002 (first entry)

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Query Match Best Local Similarity Matches 10; Conserv

Berlin K;

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07-APR-2000; 2000DE-01019173.
                                                                   (EPIG-) EPIGENOMICS AG.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99999, ABF00010-ABH99999 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but typo.int/pub/published_pct_sequences
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                                           SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Oligonucleotide SEQ ID NO 185686 for detecting SNP TSC0045759.
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Pred. No. 1.3e+03;
1; Mismatches 2; Indels
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Best Local Similarity 76.9
Matches 10, Conservative
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06-APR-2001; 2001WO-IB000713

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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                        set or oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                                                                                                             Claim 1; SEQ ID NO 210787; 29pp + Sequence Listing; German.
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WPI; 2001-657177/75
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Best Local Similarity
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ABH36074
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XW SNP; sir
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABH99998 and ABI00010-ABI82073 trepresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                             This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal; respiratory, certical nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; SS; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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designed to detect single-nucleotide polymorphisms and cytosine
methylation status.
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Claim 1; SEQ ID NO 236051; 29pp + Sequence Listing; German.
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Pred. No. 1.3e+03;
0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                   SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                              Score 9.4; DB 1; Length 13; Pred. No. 1.3e+03; 0; Mismatches 1; Indels
Sequence 13 BP; 1 A; 1 C; 5 G; 6 T; 0 U; 0 Other;
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Matches 10; Conservative
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Homo sapiens.

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                                                                                                                                   SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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designed to detect single-nucleotide polymorphisms and cytosine
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                                                                                                  Oligonucleotide SEQ ID NO 213444 for detecting SNP TSC0008090.
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12.9%; Score 9.4; DB 1; Length 13;
Best Local Similarity 90.9%; Pred. No. 1.3e+03;
Matches 10; Conservative 0; Mismatches 1; Indels
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ABH13467 standard; DNA; 13
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC09989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at
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                                                                                                                                                                                                                                                                                                                                           Claim 1; SEQ ID NO 191688; 29pp + Sequence Listing; German
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ABH44396/c
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

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               methylation status.
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Pred. No. 1.3e+03;
0; Mismatches 1; Indels
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Best Local Similarity 90.9
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                                                                         of oligonucleotides,
WPI; 2001-657177/75
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Oligonucleotide SEQ ID NO 246202 for detecting SNP TSC0060161.
ABH46225 standard; DNA; 13 BP
                                                                                                                            (first entry)
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

06-APR-2001; 2001WO-IB000713

of oligonuclectides, useful for diagnosis and cell typing, igned to detect single-nuclectide polymorphisms and cytosine WPI; 2001-657177/75.

Berlin K;

This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic diseorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99889, ABF00010-ABF99989, ABH0010-ABF99989 and ABI0010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at

12.9%; Score 9.4; DB 1; Length 13;

Query Match

Sequence 13 BP; 4 A; 0 C; 3 G; 6 T; 0 U; 0 Other;

Claim 1; SEQ ID NO 246202; 29pp + Sequence Listing; German.

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The

ö oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The coligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF9989, ABH00010-ABH99899 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic. Gaps typing, i cytosine . 0 Oligonucleotide SEQ ID NO 248177 for detecting SNP TSC0060647. Claim 1; SEQ ID NO 248177; 29pp + Sequence Listing; German. 12.9%; Score 9.4; DB 1; Length 13; 76.9%; Pred. No. 1.3e+03; tive 1; Mismatches 2; Indels designed to detect single-nucleotide polymorphisms and methylation status. Seguence 13 BP; 9 A; 2 C; 0 G; 1 T; 0 U; 1 Other; ftp.wipo.int/pub/published_pct_sequences Berlin K; BP. 06-APR-2001; 2001WO-IB000713. 07-APR-2000; 2000DE-01019173 ABH48200 standard; DNA; 13 910 TICTITGGICTT 922 (first entry) Conservative || |||||| ||: 13 TTTTTGGTATTY 1 Piepenbrock C, (EPIG-) EPIGENOMICS AG WPI; 2001-657177/75. Best Local Similarity Matches 10; Conserv WO200177384-A2 Homo sapiens. 22-FEB-2002 18-OCT-2001. ABH48200; Query Match olek A, RESULT 2295 ABH48200
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22-FEB-2002 (first entry)
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                                                                                                                                                                                                                     SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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 Pred. No. 1.3e+03;
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Best Local Similarity 90.9%;
Matches 10; Conservative
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Best Local Similarity 76.9
Matches 10; Conservative
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically prereated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99889, ABF00010-ABF99989, ABF00010-ABF99989, ABF00010-ABF99989 and ABI00010-ABF32073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WFPO at
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                                                     SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Oligonucleotide SEQ ID NO 264169 for detecting SNP TSC0064011.
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detect single-nucleotide polymorphisms and cytosine status.
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                                                                                                                                                                                     Claim 1; SEQ ID NO 95545; 29pp + Sequence Listing; German.
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ABC71543 standard; DNA; 13
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represent the oligomers described in the invention. NOTE: The sequence data for this parent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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Pred. No. 1.3e+03;
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABE99989, ABH00010-ABH99989 and ABI00010-ABI82073 data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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Oligonucleotide SEQ ID NO 71560 for detecting SNP TSC0018522.
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                                                        SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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to detect single-nucleotide polymorphisms and cytosine
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, azdiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI32073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at fitte.wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                                                                                                   Claim 1; SEQ ID NO 51053; 29pp + Sequence Listing; German.
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                                                                   Berlin K;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 12.9
Best Local Similarity 90.9
Matches 10; Conservative
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(EPIG-) EPIGENOMICS AG
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Claim 1; SEQ ID NO 51433; 29pp + Sequence Listing; German

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, contral nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form at of the printed specification, but the was obtained in electronic format from WIPO at 22222222222222222

Sequence 13 BP; 4 A; 0 C; 3 G; 6 T; 0 U; 0 Other;

12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03; ive 0; Mismatches 1; Indels Best Local Similarity 90.9%; Matches 10; Conservative 949 TTAATGTATCG 959 Query Match g ઠ

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ABF02160 standard; DNA; 13 ABF02160; RESULT 2306 ABF02160

21-FEB-2002

Oligonucleotide SEQ ID NO 102157 for detecting SNP TSC0025451.

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Homo sapiens,

WO200177384-A2

06-APR-2001; 2001WO-IB000713,

07-APR-2000; 2000DE-01019173

(EPIG-) EPIGENOMICS AG

Berlin K; Olek A, Piepenbrock C,

WPI; 2001-657177/75

Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.

Claim 1; SEQ ID NO 102157; 29pp + Sequence Listing; German.

This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) eligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretracted genemic DNA. The eligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The eligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at

Gaps ; 12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03; ive 0; Mismatches 1; Indels Sequence 13 BP; 3 A; 0 C; 2 G; 8 T; 0 U; 0 Other; Conservative Query Match Best Local Similarity Matches 10; Conserv

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947 GITTAATGTAT 957 11 GTTTATTGTAT . 요

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ABC03281 standard; DNA; 13 BP.

ABC03281;

20-FEB-2002

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Gaps

0

Oligonucleotide SEQ ID NO 3272 for detecting SNP TSC0001238.

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Homo sapiens.

WO200177384-A2.

18-OCT-2001

06-APR-2001; 2001WO-IB000713

07-APR-2000; 2000DE-01019173 (EPIG-) EPIGENOMICS Berlin K; Olek A, Piepenbrock C,

WPI; 2001-65717/75

Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.

Claim 1; SEO ID NO 3272; 29pp + Sequence Listing; German.

This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABE99889, ABF00010-ABE99889, ABF00010-ABE99889, ABF00010-ABE99899, ABF00010-ABE99899 and ABI00010-ABI32073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WFPO at the printed specification, but ftp.wipo.int/pub/published_pot_sequences

Sequence 13 BP; 2 A; 6 C; 0 G; 4 T; 0 U; 1 Other;

ö Length 13; Score 9.4; DB 1; Length 13 Pred. No. 1.3e+03; 0; Mismatches 1; Indels 12.9%; Query Match
12.9
Best Local Similarity 90.9
Matches 10; Conservative

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Gaps

930 ATCCCTCCTCT 940

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RESULT 2308 ABC03282/c

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range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC09989, ABF00010-ABF9988, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence was obtained in electronic format from WIPO at the printed specification, but the wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a
                                                                                                                                                                                                                                                                                                                                                                                                                                                              This invention describes novel oligonucleotide primers or peptide nucleic
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                                                                                                                                                                                                                                                                                                                          Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine
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Pred. No. 1.3e+03;
0; Mismatches 1; Indels
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90.9%;
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les 10; Conservative
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                                                                                                                                                                                                                                                                                                                                                                           methylation status.
                       WO200177384-A2.
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                                                                                                                                                                           SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                   Oligonucleotide SEQ ID NO 3273 for detecting SNP TSC0001238.
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Pred. No. 1.3e+03;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1; SEQ ID NO 3273; 29pp + Sequence Listing; German.
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    ABC03282 standard; DNA; 13
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                                                                                       20-FEB-2002 (first entry)
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olek A,

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Query Match

Matches

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Gaps

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PMA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99899 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at of oligonucleotides, useful for diagnosis and cell typing, igned to detect single-nucleotide polymorphisms and cytosine Claim 1; SEQ ID NO 104550; 29pp + Sequence Listing; German. WPI; 2001-657177/75 designed to detect methylation status. Set

Sequence 13 BP; 8 A; 3 C; 1 G; 0 T; 0 U; 1 Other;

ö Length 13; 2; Indels Score 9.4; DB 1; Pred. No. 1.3e+03; 1; Mismatches 2; 12.9%; 10; Conservative Query Match Best Local Similarity Matches

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ABC55216 standard; DNA; 13

(first entry) 21-FEB-2002

ABC55216;

Oligonucleotide SEQ ID NO 55233 for detecting SNP TSC0015098.

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Homo sapiens

WO200177384-A2

18-OCT-2001

06-APR-2001; 2001WO-IB000713.

07-APR-2000; 2000DE-01019173;

(EPIG-) EPIGENOMICS AG

Berlin K; ບັ Piepenbrock WPI; 2001-657177/75. olek A,

ligonucleotides, useful for diagnosis and cell typing, it o detect single-nucleotide polymorphisms and cytosine of oligonucleotides, designed

methylation status.

Claim 1; SEQ ID NO 55233; 29pp + Sequence Listing; German.

This invention describes novel oligonucleotide primers or peptide inucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory,

central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABCO0010-ABC09989, ABF00010-ABF99989, ABF00010-ABF99989, ABF00010-ABF99989, ABF00010-ABF99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence was obtained in electronic format from WIPO at thick parinted specification, but ftp.wipo.int/pub/published_pct_sequence 888888888888

Sequence 13 BP; 4 A; 0 C; 3 G; 6 T; 0 U; 0 Other;

Gaps ; 0 Query Match 12.9%; Score 9.4; DB 1; Length 13; Best Local Similarity 90.9%; Pred. No. 1.3e+03; Matches 10; Conservative 0; Mismatches 1; Indels

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946 GGTTTAATGTA 956 GGTTTTATGTA 11 Н

셤 8

RESULT 2312

BP. ABC32309 standard; DNA; 13 ABC32309,

ABC32309;

(first entry) 20-FEB-2002

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Gaps

Oligonucleotide SEQ ID NO 32326 for detecting SNP TSC0010079.

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Homo sapiens

WO200177384-A2.

18-OCT-2001

06-APR-2001; 2001WO-IB000713

07-APR-2000; 2000DE-01019173

(EPIG-) EPIGENOMICS

Berlin Olek A, Piepenbrock C,

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WPI; 2001-657177/75.

octor congonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.

Claim 1; SEQ ID NO 32326; 29pp + Sequence Listing; German.

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences

Sequence 13 BP; 9 A; 3 C; 0 G; 1 T; 0 U; 0 Other;

ö Gaps .. 0 Query Match 12.9%; Score 9.4; DB 1; Length 13; Best Local Similarity 90.9%; Pred. No. 1.3e+03; Matches 10; Conservative 0; Mismatches 1; Indels

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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99889, ABF00010-ABF99899, ABH00010-ABF99899 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic formmat from WIPO at
                                                       SNP; single nuclectide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                Oligonucleotide SEQ ID NO 85638 for detecting SNP TSC0021525.
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ABC85621 standard; DNA; 13
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Best Local Similarity
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
Oligonucleotide SEQ ID NO 111070 for detecting SNP TSC0027729.
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RESULT 23
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ABC86657 standard; DNA; 13 BP.
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This invention describes novel oligonucleotide primers or peptide nucleic "acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                  Claim 1; SEQ ID NO 86674; 29pp + Sequence Listing; German.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretraeted genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular, and metabolic discorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABC0010-ABE99989, ABC0010-ABE99989, ABC0010-ABE99989, ABC0010-ABE99989 and ABI00010-ABE82073 represent the oligomers described in the invention. NOTE: The sequence was obtained in electronic formmat from WIPO at the printed specification, but the wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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90.9%;
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                                                               ABC64623 standard; DNA; 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                         Score 9.4; DB 1; Length 13;
Pred. No. 1.3e+03;
0; Mismatches 1; Indels
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                                                                    Sequence 13 BP; 2 A; 6 C; 1 G; 4 T; 0 U; 0 Other;
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was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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WO200177384-A2 Homo sapiens

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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 trepresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                oer or oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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central nervous system; gastrointestinal; respiratory; immune; metabolic.
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and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABE99989, ABF0010-ABE99989, ABF0010-ABE99989, ABF0010-ABE99989 and ABI0010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but two obtained in electronic format from WIPO at
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designed to detect single-nucleotide polymorphisms and cytosine
methylation status.
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Pred. No. 1.3e+03;
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                                                                                                                                                                                                                                                                                                        SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Pred. No. 1.3e+03;
1; Mismatches 2; Indels
Score 9.4; DB 1; Length 13;
Pred. No. 1.30+03;
1; Mismatches 2; Indels
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76.9%;
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Best Local Similarity 76.9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SNP, single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                  SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic
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                                                                  Oligonucleotide SEQ ID NO 133100 for detecting SNP TSC0033208.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a renge of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                      This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABH32073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at fixe wipo int/pub/published_pct_sequences
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oligonuclectides, useful for diagnosis and cell typing, is ed to detect single-nuclectide polymorphisms and cytosine
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                                                             Claim 1; SEQ ID NO 218936; 29pp + Sequence Listing; German.
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Pred. No. 1.3e+03;
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Matches 10; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
              represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073
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designed to detect single-nucleotide polymorphisms and cytosine
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                                                                                                                                            Score 9.4; DB 1; Length 13; Pred. No. 1.3e+03; 0; Mismatches 1; Indels
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Pred. No. 1.3e+03;
0; Mismatches 1; Indels
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                                                                                                            Sequence 13 BP; 7 A; 0 C; 1 G; 5 T; 0 U; 0 Other;
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ftp.wipo.int/pub/published_pct_sequences
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methylation status.
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Best Local Similarity
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABCO0010

Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine

methylation status.

Claim 1; SEQ ID NO 194179; 29pp + Sequence Listing; German.

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Query Match
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                                                                                                                                                                     SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                  Oligonucleotide SEQ ID NO 197817 for detecting SNP TSC0048685.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1; SEQ ID NO 197817; 29pp + Sequence Listing; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03; tive 0; Mismatches 1; Indels
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABR00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 the preparent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99899, ABF00010-ABE99899, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; 88; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                             Claim 1; SEQ ID NO 199387; 29pp + Sequence Listing; German.
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Claim 1; SEQ ID NO 176915; 29pp + Sequence Listing; German.

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            This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at first pub/published_pct_sequences
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABE9989, ABF00010-ABE9989, ABH00010-ABE9989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic formmat from WIPO at
                                                                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                          Oligonucleotide SEQ ID NO 227834 for detecting SNP TSC0055556.
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                    ABH27857 standard; DNA; 13 BP.
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RESULT 23
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                                                                                                                                                                                                                                                                                                                                                                                           SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                   Score 9.4; DB 1; Length 13;
Pred. No. 1.3e+03;
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                 Sequence 13 BP; 2 A; 0 C; 2 G; 9 T; 0 U; 0 Other;
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a
                                                                                                                                                                      This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metholic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABE99989, ABF00010-ABF99989, ABH00010-ABH99999 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                  ides, useful for diagnosis and cell typing, is single-nucleotide polymorphisms and cytosine
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                                                                                                                                     Claim 1; SEQ ID NO 204874; 29pp + Sequence Listing; German.
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                                                                                                    methylation status.
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range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF9989, ABH00010-ABH99899 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form par of the printed specification, but was obtained in electronic format from WIPO at fire wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABE99989, ABF00010-ABE99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                    SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Length 13;

12.9%; Score 9.4; DB 1; 90.9%; Pred. No. 1.3e+03;

Query Match Best Local Similarity

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99889, ABF00010-ABF99889, ABH00010-ABH99889 and ABT00010-ABI82073 trepresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from NIPO at
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                                                                   Oligonucleotide SEQ ID NO 158531 for detecting SNP TSC0039907.
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                                                               This invention describes novel oligonucleotide primers or peptide nucleic acid (PMA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABE99989, ABF00010-ABE99899, ABF00010-ABE99899, ABF00010-ABE99899 and ABI00010-ABE82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but fip.wipo.int/pub/published_pct_sequences
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                                  Claim 1; SEQ ID NO 160973; 29pp + Sequence Listing; German.
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                                                                                                                                  SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                             Oligonucleotide SEQ ID NO 214460 for detecting SNP TSC0052165.
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peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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ô This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99899, ABF00010-ABF99899 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but fur wipo.int/pub/published_pct_sequences SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic. Gaps set or oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status. Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status. . 0 Oligonucleotide SEQ ID NO 245826 for detecting SNP TSC0010699, Claim 1; SEQ ID NO 245825; 29pp + Sequence Listing; German. Score 9.4; DB 1; Length 13; Pred. No. 1.3e+03; 0; Mismatches 1; Indels Sequence 13 BP; 3 A; 0 C; 1 G; 8 T; 0 U; 1 Other; 꿌 Berlin K; Berlin ABH45849 standard; DNA; 13 BP. 12.9%; 90.9%; 06-APR-2001; 2001WO-IB000713. 07-APR-2000; 2000DE-01019173 22-FEB-2002 (first entry) Query Match 12.9 Best Local Similarity 90.9 Matches 10, Conservative 943 ATTGGTTTAAT 953 Olek A, Piepenbrock C, 1 Arrerrraar 11 (EPIG-) EPIGENOMICS AG. WPI; 2001-657177/75. WPI; 2001-657177/75. WO200177384-A2. Homo sapiens. 18-OCT-2001 ABH45849; olek A, RESULT 2350

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Oligonucleotide SEQ ID NO 247682 for detecting SNP TSC0060535.

22-FEB-2002 (first entry)

ABH47705 standard; DNA; 13 BP

acid (PNA) oligomers for detecting single nuclectide polymorphisms (SNP) and cytosine methylation status in chemically precreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99899, ABC0010-ABE99899 ABH0010-ABH99989 and ABI0010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at the printed specification, but ftp.wipo.int/pub/published_pct_sequences

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Matches 10; Conservative

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disconders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo.int/pub/published_pct_sequences SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic. Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine methylation status. Claim 1; SEQ ID NO 247682; 29pp + Sequence Listing; German. Sequence 13 BP; 6 A; 3 C; 0 G; 4 T; 0 U; 0 Other; Berlin K; 06-APR-2001; 2001WO-IB000713. 07-APR-2000; 2000DE-01019173 Olek A, Piepenbrock C, (EPIG-) EPIGENOMICS AG. WPI; 2001-657177/75. WO200177384-A2. Homo sapiens 18-OCT-2001.

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This invention describes novel oligonucleotide primers or peptide nucleic

Claim 1; SEQ ID NO 245826; 29pp + Sequence Listing; German.

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                 SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                               Oligonucleotide SEQ ID NO 264228 for detecting SNP TSC0064030.
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(first entry)
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Homo sapiens

RESULT 2353 ABH64251/c ID ABH64251 standard; DNA; 13 BP.

This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretracted genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but thous, was obtained in electronic format from WIPO at Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status. Claim 1; SEQ ID NO 265110; 29pp + Sequence Listing; German. Sequence 13 BP; 1 A; 5 C; 0 G; 6 T; 0 U; 1 Other; 06-APR-2001; 2001WO-IB000713. 07-APR-2000; 2000DE-01019173, Olek A, Piepenbrock C, (EPIG-) EPIGENOMICS AG WPI; 2001-657177/75 WO200177384-A2 18-OCT-2001.

Query Match 12.9%; Score 9.4; DB 1; Length 13; Best Local Similarity 90.9%; Pred. No. 1.3e+03; Matches 10; Conservative 0; Mismatches 1; Indels

935 TCCTCTTCATT 945 3 recreirecti 13 à

ABH65663 standard; DNA; 13 BP. ABH65663; RESULT 233
ABH65663/A
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AC ABH6
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(first entry) 22-FEB-2002

Oligonucleotide SEQ ID NO 265640 for detecting SNP TSC0064381.

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic. SNP; sin

Homo sapiens

WO200177384-A2

18-OCT-2001

06-APR-2001; 2001WO-IB000713

07-APR-2000; 2000DE-01019173.

(EPIG-) EPIGENOMICS AG

Berlin K; Piepenbrock C, olek A,

WPI; 2001-657177/75

Set of oligonuclectides, useful for diagnosis and cell typing, i designed to detect single-nuclectide polymorphisms and cytosine methylation status.

Claim 1; SEQ ID NO 265640; 29pp + Sequence Listing; German.

This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at

Sequence 13 BP; 7 A; 2 C; 0 G; 4 T; 0 U; 0 Other;

Gaps 6 Query Match 12.9%; Score 9.4; DB 1; Length 13; Best Local Similarity 90.9%; Pred. No. 1.3e+03; Matches 10; Conservative 0; Mismatches 1; Indels

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ò 셤 RESULT 2356

ABC93439 standard; DNA; 13

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ABC93439;

(first entry) 21-FEB-2002

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Gaps

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Oligonucleotide SEQ ID NO 93456 for detecting SNP TSC0023347.

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Homo sapiens.

WO200177384-A2.

18-OCT-2001,

06-APR-2001; 2001WO-IB000713.

07-APR-2000; 2000DE-01019173.

(EPIG-) EPIGENOMICS

Berlin K; Olek A, Piepenbrock C,

WPI; 2001-657177/75

Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.

Claim 1; SEQ ID NO 93456; 29pp + Sequence Listing; German.

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99889, ABF00010-ABF99899, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                     SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                                                          Oligonucleotide SEQ ID NO 71559 for detecting SNP TSC0018522.
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Best Local Similarity 90.9
Matches 10, Conservative
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  930 ATCCCTCCTCT 940
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oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99889, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NoTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at fig. wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                             12.9%; Score 9.4; DB 1; Length 13; 76.9%; Pred. No. 1.3e+03; tive 1; Mismatches 2; Indels
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nes 10; Conservative
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RESULT 2357

Matches

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Sequence 13 BP; 1 A; 0 C; 2 G; 9 T; 0 U; 1 Other;
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
Oligonucleotide SEQ ID NO 21566 for detecting SNP TSC0004330.
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Homo sapiens

MOD OCE TO T4:40:T3 7004

WO200177384-A2

18-OCT-2001

06-APR-2001; 2001WO-IB000713.

07-APR-2000; 2000DE-01019173

(EPIG-) EPIGENOMICS AG

Berlin K; Olek A, Piepenbrock C,

WPI; 2001-657177/75

Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.

Claim 1; SEQ ID NO 21566; 29pp + Sequence Listing, German.

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretraeted genemic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99999, ABF00010-ABF99999 and ABI00010-ABF3073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wibo.int/pub/published_pct_sequences

Sequence 13 BP; 0 A; 5 C; 1 G; 7 T; 0 U; 0 Other;

0; Gaps 12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03; ive 0; Mismatches 1; Indels Local Similarity 90.9 nes 10; Conservative Query Match

912 CTTTGGTCTTT 922 |||| |||||| 2 CTTTCGTCTTT 12

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RESULT 2360

ABC71615 standard; DNA; 13 BP.

ABC71615;

21-FEB-2002 (first entry)

Oligonucleotide SEQ ID NO 71632 for detecting SNP TSC0018535.

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Homo sapiens

WO200177384-A2

18-OCT-2001.

06-APR-2001; 2001WO-IB000713

07-APR-2000; 2000DE-01019173

(EPIG-) EPIGENOMICS AG

Berlin K; Piepenbrock C, olek A,

WPI; 2001-657177/75

Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.

Claim 1; SEQ ID NO 71632; 29pp + Sequence Listing; German.

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF9989, ABH0010-ABH99899 and ABI0010-ABI82073 trepresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo.int/pub/published_pot_sequences

Sequence 13 BP; 8 A; 2 C; 0 G; 2 T; 0 U; 1 Other;

Gaps .; 0 Query Match
12.9%; Score 9.4; DB 1; Length 13;
Best Local Similarity 76.9%; Pred. No. 1.3e+03;
Matches 10; Conservative 1; Mismatches 2; Indels

ò 셤 2361 ABC97969 RESULT

ABC97969 standard; DNA; 13 BP.

ABC97969;

21-FEB-2002 (first entry)

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Oligonucleotide SEQ ID NO 97986 for detecting SNP TSC0024337.

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Homo sapiens.

WO200177384-A2.

18-OCT-2001.

06-APR-2001; 2001WO-IB000713.

07-APR-2000; 2000DE-01019173

(EPIG-) EPIGENOMICS AG.

Olek A, Piepenbrock C,

Berlin K;

WPI; 2001-657177/75.

Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.

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Matches
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                        This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99899, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from MIPO at
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Claim 1; SEQ ID NO 97986; 29pp + Sequence Listing; German.
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                                                                                                                                                                                                                                           Sequence 13 BP; 3 A; 1 C; 0 G; 9 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                           0; Mismatches
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                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                       Score 9.4; DB 1; Length 13;
Pred. No. 1.3e+03;
1; Mismatches 2; Indels
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                                                  Sequence 13 BP; 2 A; 0 C; 4 G; 6 T; 0 U; 1 Other;
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Matches 10; Conservative
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les 10; Conserv
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Berlin K;

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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABF32073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                 Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 1; SEQ ID NO 3271; 29pp + Sequence Listing; German.
                                                                                                                                      36-APR-2001; 2001WO-IB000713
                                                                                                                                                                          07-APR-2000; 2000DE-01019173
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                                                                                                                                                                                          SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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designed to detect single-nuclectide polymorphisms and cytosine
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                                   ABC75194 standard; DNA; 13
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Best Local Similarity 90.9
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11 CTTTTTCCCT 1
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RESULT 2364
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                                                      12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03; ive 0; Mismatches 1; Indels
Sequence 13 BP; 4 A; 0 C; 6 G; 2 T; 0 U; 1 Other;
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oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The
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designed to detect single-nucleotide polymorphisms and cytosine
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Piepenbrock C,
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Sequence 13 BP; 2 A; 0 C; 2 G; 8 T; 0 U; 1 Other;

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Oligonucleotide SEQ ID NO 106772 for detecting SNP TSC0026727.

(first entry)

21-FEB-2002

ABF06775;

ABF06775 standard; DNA; 13 BP.

RESULT 2368

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disonders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99999, ABF00010-ABF99999, ABH00010-ABF99999 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from NIPO at
                                                                                                                                             SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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cytosine
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    methylation status.
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                                                                                                                                                                                                                        Homo sapiens
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Berlin K;

12.9%; Score 9.4; DB 1; Length 13;

Query Match

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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                         Oligonucleotide SEQ ID NO 8873 for detecting SNP TSC0002401.
               20-FEB-2002 (first entry)
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                                                                                                                                                                                                                                         SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                  Oligonucleotide SEQ ID NO 58002 for detecting SNP TSC0015581.
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                 1; Indels
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  90.9%; Pred. No. 1.3e+03;
               Mismatches
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ABC57985 standard; DNA; 13 BP.
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Best Local Similarity
Matches 10; Conserva
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ABC57985/c
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                                                   WO200177384-A2.
Homo sapiens.
                                                                                                             18-OCT-2001.
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ABC08882 standard; DNA; 13 BP.

RESULT 2370 ABC08882/c ID ABC08 XX AC ABC08

ABC08882;

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and methololic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99988 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                                                                                                                                                                                                                                                                                                 Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine
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Best Local Similarity
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ABC36332
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AC ABC36333
AC ABC36333
DT 20-FEB-2
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KW SNP; SI
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Set of oligonucleotides, useful for diagnosis and cell typing, is

Berlin K;

Piepenbrock C,

olek A,

WPI; 2001-657177/75

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989 and ABI00010-ABF82073
                                                                                                             This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99899, ABF00010-ABF99899, ABH00010-ABH99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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Pred. No. 1.3e+03;
1; Mismatches 2; Indels
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                                                                             Claim 1; SEQ ID NO 36349; 29pp + Sequence Listing; German.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           1; Mismatches
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Best Local Similarity
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DNA; 13 BP.

(first entry)

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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                  Oligonucleotide SEQ ID NO 14564 for detecting SNP TSC0003286.
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                                                                       ABC14557 standard;
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represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                               SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                       Gaps
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12.9%; Score 9.4; DB 1; Length 13;
Best Local Similarity 90.9%; Pred. No. 1.3e+03;
Matches 10; Conservative 0; Mismatches 1; Indels
                                                                                                      Query Match 12.9%; Score 9.4; DB 1; Length 13; Best Local Similarity 76.9%; Pred. No. 1.38+03; Matches 10; Conservative 1; Mismatches 2; Indels
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                                                                         Sequence 13 BP; 5 A; 3 C; 0 G; 4 T; 0 U; 1 Other;
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                                                                                                                                                                                                                                                                          ABC88572 standard; DNA; 13
                                                                                                                                                                  948 TTTAATGTATCGC 960
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Berlin K;

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABF3073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the pub/published_pct_sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03; tive 0; Mismatches 1; Indels
Claim 1; SEQ ID NO 14564; 29pp + Sequence Listing; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 13 BP; 0 A; 9 C; 0 G; 4 T; 0 U; 0 Other;
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hes 10; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                                                                                                                                 Claim 1; SEQ ID NO 63731; 29pp + Sequence Listing; German.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99999, ABF00010-ABF99999, ABH00010-ABH99999 and ABI00010-ABF3073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic formmat from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                              Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                Berlin K;
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nes 10; Conservative
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                                               Piepenbrock C,
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                (EPIG-) EPIGENOMICS AG.
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                                                olek A,
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Sequence 13 BP; 3 A; 0 C; 4 G; 6 T; 0 U; 0 Other;

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nuclectide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, contral nervous system, cardiovaecular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABH99989 and ABI00010-ABH82073 tepresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at 888888888888888888

Sequence 13 BP; 2 A; 0 C; 2 G; 9 T; 0 U; 0 Other;

Gaps . 12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03; ive 0; Mismatches 1; Indels Best Local Similarity 90.5 Matches 10; Conservative Query Match

947 GITTAATGTAT 957 Grrraargrrr 12

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ABF26356 standard; DNA; 13 RESULT 2379

21-FEB-2002 ABF26356;

BP.

Oligonucleotide SEQ ID NO 126353 for detecting SNP TSC0031615.

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Berlin K;

Olek A, Piepenbrock C,

(EPIG-) EPIGENOMICS AG.

06-APR-2001; 2001WO-IB000713. 07-APR-2000; 2000DE-01019173

WO200177384-A2

18-OCT-2001.

Homo sapiens.

WO200177384-A2

18-OCT-2001

06-APR-2001; 2001WO-IB000713.

07-APR-2000; 2000DE-01019173

(EPIG-) EPIGENOMICS AG.

Berlin K; Olek A, Piepenbrock C,

WPI; 2001-657177/75.

Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.

Claim 1; SEQ ID NO 126353; 29pp + Sequence Listing; German.

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF9989, ABF00010-ABF9989 and ABI00010-ABF3073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo.int/pub/published_pct_sequences

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                  12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03; ive 0; Mismatches 1; Indels
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Query Match
Best Local Similarity 90.2
These 10; Conservative
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ID ABF3
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99889, ABF00010-ABF99989, ABH00010-ABF99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status. Claim 1; SEQ ID NO 131635; 29pp + Sequence Listing; German WPI; 2001-657177/75.

Gaps .. 12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03; ive 0; Mismatches 1; Indels Sequence 13 BP; 1 A; 0 C; 3 G; 9 T; 0 U; 0 Other; Query Match
Best Local Similarity 90.9
Matches 10; Conservative

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RESULT 2381 ABF31787/c

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WO200177384-A2
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                                                                                                                                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                        Oligonucleotide SEQ ID NO 131784 for detecting SNP TSC0032896,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 1; SEQ ID NO 131784; 29pp + Sequence Listing; German
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Pred. No. 1.3e+03;
0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             was obtained in electronic format from W1 ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                    (first entry)
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   DNA;
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   ABF31787 standard;
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABE09989, ABF00010-ABE99989, ABH00010-ABE99989, and ABI00010-ABE82073 capresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic formmat from WIPO at
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 1; SEQ ID NO 136949; 29pp + Sequence Listing; German.
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06-APR-2001; 2001WO-IB000713.
                                                              07-APR-2000; 2000DE-01019173
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABH99989 and ABI00010-ABH92073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but typo.wipo.int/pub/published_pct_sequences
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                                   Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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designed to detect single-nucleotide polymorphisms and cytosine
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                                                                                                         Claim 1; SEQ ID NO 218258; 29pp + Sequence Listing; German.
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Pred. No. 1.3e+03;
0; Mismatches 1; Indels
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90.9%;
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WPI; 2001-657177/75
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central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABCO010 -ABC09989, ABF00010-ABF9989, ABF00010-ABF9989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at thichpub/published_pct_sequences
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                                                                                                                                                      Sequence 13 BP; 2 A; 6 C; 0 G; 5 T; 0 U; 0 Other;
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Matches 10; Conserv
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Oligonucleotide SEQ ID NO 148523 for detecting SNP TSC0037485.
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정, 원
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                                                                                                                                                                                                                                                                                                                                                                                                         SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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ABF48526 standard; DNA; 13
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Matches 10; Conservative
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; SEQ ID NO 148523; 29pp + Sequence Listing; German.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ftp.wipo.int/pub/published_pct_sequences
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es 10; Conservative
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schultz1-899.rng

Claim 1; SEQ ID NO 226134; 29pp + Sequence Listing; German.

This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABE99989, ABF00010-ABE99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo.int/pub/published_pct_sequence

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Query Match
Best Local Similarity 90.9
Matches 10; Conservative
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                                                                                                                                                                                                                                                     This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABH3073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                          Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                                                                                                                                            Claim 1; SEQ ID NO 199596; 29pp + Sequence Listing; German.
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                                                                                              Berlin K;
06-APR-2001; 2001WO-IB000713.
                              07-APR-2000; 2000DE-01019173
                                                                                            Olek A, Piepenbrock C,
                                                             (EPIG-) EPIGENOMICS AG
                                                                                                                           WPI; 2001-657177/75.
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Gaps

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12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03; ive 0; Mismatches 1; Indels

924 CCTTTTATCCC 934 CCATTTATCCC 11 ABH02534 standard; DNA; 13 BP.

Sequence 13 BP; 2 A; 6 C; 0 G; 5 T; 0 U; 0 Other;

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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                     Oligonucleotide SEQ ID NO 202511 for detecting SNP TSC0049776.
                                              22-FEB-2002 (first entry)
                      ABH02534;
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                                                                                                                                                                                                                                                SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                          Oligonucleotide SEQ ID NO 226134 for detecting SNP TSC0055119.
                    12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03; ive 0; Mismatches 1; Indels
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Best Local Similarity 90.9
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ABH26157

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Claim 1; SEQ ID NO 202511; 29pp + Sequence Listing; German. WPI; 2001-657177/75. Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.

Berlin K;

Olek A, Piepenbrock C, (EPIG-) EPIGENOMICS AG.

WPI; 2001-657177/75.

06-APR-2001; 2001WO-IB000713. 07-APR-2000; 2000DE-01019173.

WO200177384-A2 Homo sapiens,

18-OCT-2001.

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PMA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastroinfestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010 ABC0999, ABC090010-ABC99989, ABC0010-ABC9988, ABC0010-ABC9988, ABC0010-ABC9988, ABC0010-ABC9988, ABC0010-ABC9988, ABC0010-ABC9988, ABC0010-ABC9988, ABC0010-ABC9988, ABC0010-ABC9988, ABC0010-ABC988, ABC0010-ABC9988, ABC00010-ABC9988, ABC0010-ABC9988, ABC0010-ABC9
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Homo sapiens.
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ABH37215;
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           RESULT 2392
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                                                                                                                                                                                                                                                                                             SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                      Gaps
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Pred. No. 1.3e+03;
0; Mismatches 1; Indels
                                                           Score 9.4; DB 1; Length 13; Pred. No. 1.3e+03;
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                                    Seguence 13 BP; 6 A; 0 C; 6 G; 1 T; 0 U; 0 Other;
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was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                  0; Mismatches
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90.9%;
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                                                                        Local Similarity 90.9
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ses 10; Conservative
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                                                                                                                                                                                                                                                                                                                                                   Homo sapiens
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                                                                                                                                                                                                                                         SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                   Oligonucleotide SEQ ID NO 184013 for detecting SNP TSC0045426.
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BP.
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ABF84016 standard; DNA; 13
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central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                                           Olek A, Piepenbrock C,
                                                                                                                                                                                                                                                              (EPIG-) EPIGENOMICS AG
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                                                                                                                                                                                                                                                                                                                                                     WPI; 2001-657177/75.
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                                                                                    WO200177384-A2
                                            Homo sapiens.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABE99989, ABF00010-ABE99989, ABF00010-ABE99989, ABF00010-ABE99989 and ABI00010-ABE92073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                                                                                                                                                                                                                              Claim 1; SEQ ID NO 163908; 29pp + Sequence Listing; German.
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Berlin K;
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Piepenbrock C,
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and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for disquosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABC0010-ABF9989, ABH00010-ABF9989, ABH00010-ABF9989, ABH00010-ABF9989, ABF00010-ABF9989, ABF00010010-ABF9989, ABF00010-ABF9989, ABF9989, ABF9989
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Best Local Similarity
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                                                                                                                                           SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                        Oligonucleotide SEQ ID NO 246201 for detecting SNP TSC0060161.
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76.9%;
ABH46224 standard; DNA; 13
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Sequence 13 BP; 5 A; 0 C; 8 G; 0 T; 0 U; 0 Other;

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 targresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from NIPO at
                                                                                                   SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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designed to detect single-nucleotide polymorphisms and cytosine
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                                                                   Oligonucleotide SEQ ID NO 251279 for detecting SNP TSC0061339.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                was obtained in electronic format from Wiftp.wipo.int/pub/published_pct_sequences
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABE09989, ABF00010-ABE99899, ABF00010-ABE99989, ABF00010-ABE99989, ABF00010-ABE99989 and ABI00010-ABE82073 tepsement the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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ABH57301 standard; DNA; 13 BP.
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ligonucleotides, useful for diagnosis and cell typing, is to detect single-nucleotide polymorphisms and cytosine

Set of oligonucleotides,

methylation status.

Berlin K;

Olek A, Piepenbrock C, WPI; 2001-657177/75.

(EPIG-) EPIGENOMICS

07-APR-2000; 2000DE-01019173

06-APR-2001; 2001WO-IB000713

18-OCT-2001

ABH51302;

Claim 1; SEQ ID NO 253876; 29pp + Sequence Listing; German.

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                                                                                                                 This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99899, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine
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designed to detect single-nucleotide polymorphisms and cytosine
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                                                                                Claim 1; SEQ ID NO 257278; 29pp + Sequence Listing; German
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ftp.wipo.int/pub/published_pct_sequences
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Matches 10; Conservative
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13 TGTTTTATTGTAY 1
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                                         methylation status.
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-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at the print pub/published_pot_sequences
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                                                                                                            Seguence 13 BP; 4 A; 0 C; 2 G; 7 T; 0 U; 0 Other;
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Best Local Similarity
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Best Local Similarity
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Page 1047

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ABC68201 standard; DNA; 13 BP. ABC68203

ABC68201;

(first entry) 21-FEB-2002

Oligonucleotide SEQ ID NO 68218 for detecting SNP TSC0017803.

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Homo sapiens.

WO200177384-A2.

18-OCT-2001

06-APR-2001; 2001WO-IB000713,

07-APR-2000; 2000DE-01019173.

(EPIG-) EPIGENOMICS AG

Berlin K; Olek A, Piepenbrock C,

WPI; 2001-657177/75.

Set of oligonuclectides, useful for diagnosis and cell typing, is designed to detect single-nuclectide polymorphisms and cytosine methylation status.

Claim 1; SEQ ID NO 68218; 29pp + Sequence Listing; German.

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABE99989, ABF00010-ABE99899, ABF00010-ABE99899, ABF00010-ABE99899 and ABI00010-ABE3073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but ftp.wipo.int/pub/published_pct_sequences

Sequence 13 BP; 2 A; 4 C; 0 G; 7 T; 0 U; 0 Other;

12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03; ive 0; Mismatches 1; Indels Query Match
Best Local Similarity 90.9
Matches 10; Conservative

935 TCCTCTTCATT 945

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3 TCCTCTACATT 13

RESULT 2404
ABC22206
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AC ABC22201
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DT 20-FEB-,
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ABC22206 standard; DNA; 13 BP.

ABC22206;

(first entry) 20-FEB-2002 Oligonucleotide SEQ ID NO 22223 for detecting SNP TSC0004408.

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Homo sapiens

WO200177384-A2

18-OCT-2001

06-APR-2001; 2001WO-IB000713.

07-APR-2000; 2000DE-01019173.

(EPIG-) EPIGENOMICS AG

Ϋ́ Berlin Olek A, Piepenbrock C,

WPI; 2001-657177/75.

set or oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.

Claim 1; SEQ ID NO 22223; 29pp + Sequence Listing; German. .

This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABE99999, ABF00010-ABE99999, ABF00010-ABE99999, ABF00010-ABE99999 and ABI00010-ABE92073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at fire printed specification, but fip.wipo.int/pub/published_pct_sequences

Sequence 13 BP; 4 A; 0 C; 4 G; 4 T; 0 U; 1 Other;

. 0 Query Match 12.9%; Score 9.4; DB 1; Length 13; Best Local Similarity 90.9%; Pred. No. 1.3e+03; Matches 10; Conservative 0; Mismatches 1; Indels

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RESULT 2405

ABC97648 standard; DNA; 13 BP. ABC97648

ABC97648;

21-FEB-2002 (first entry)

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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic. Oligonucleotide SEQ ID NO 97665 for detecting SNP TSC0024259.

Homo sapiens.

WO200177384-A2.

18-OCT-2001.

06-APR-2001; 2001WO-IB000713

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99889, ABF00010-ABF99889, ABH00010-ABH99989 and ABI00010-ABF32073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic formmat from WIPO at
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Pred. No. 1.3e+03;
0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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designed to detect single-nucleotide polymorphisms and cytosine
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                                                                                                                                         Berlin K;
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07-APR-2000; 2000DE-01019173
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                                                                    (EPIG-) EPIGENOMICS AG
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Query Match

8 셤 ABC76020;

This invention describes novel oligomucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically prereated genomic DNA. The oligomucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic discorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABC0010-ABC99989, ABC0010-ABC99989 and ABI00010-ABC0010 card for this patent did not form part of the printed specification, but was obtained in electronic formmat from WIPO at the printed specification, but ftp.wipo.int/pub/published_pct_sequences

olek A,

(first entry)

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                              SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                          ABC51406 standard; DNA; 13 BP.
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ABC76825/
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                          12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03; rive 0; Mismatches 1; Indels
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                          Sequence 13 BP; 4 A; 0 C; 8 G; 1 T; 0 U; 0 Other;
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Berlin K;

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                                Score 9.4; DB 1; Length 13;
Pred. No. 1.3e+03;
0; Mismatches 1; Indels
Sequence 13 BP; 2 A; 0 C; 5 G; 6 T; 0 U; 0 Other;
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ilarity 90.9%;
Conservative
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                         Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                              Claim 1; SEQ ID NO 103496; 29pp + Sequence Listing; German.
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                                                                                                                                                                                                                                                                                                                                            Claim 1; SEQ ID NO 76842; 29pp + Sequence Listing; German.
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                                  WO200177384-A2
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   Homo sapiens.
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range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99899, ABF00010-ABF99989, ABF00010-ABF99989, ABF00010-ABF99989, ABF00010-ABF99989, Data for this parent did not form par to five printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardicvascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Pred. No. 1.3e+03;
1; Mismatches 2; Indels
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76.9%;
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Best Local Similarity
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                                                                                                                                                                                                                                                     SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                    Oligonucleotide SEQ ID NO 54230 for detecting SNP TSC0014889.
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ABC79371/c
ID ABC79371 standard; DNA; 13
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Length 13;

Score 9.4; DB 1; Pred. No. 1.3e+03;

12.9%;

Query Match Best Local Similarity

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 trepseent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                                       SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                       Oligonucleotide SEQ ID NO 79388 for detecting SNP TSC0020177
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21-FEB-2002 (first entry)
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ses 10; Conservative
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Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine Berlin K; Piepenbrock C, WPI; 2001-657177/75 olek A,

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Homo sapiens

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                                                    This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99889, ABF00010-ABF99899, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form par of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Oligonucleotide SEQ ID NO 11864 for detecting SNP TSC0002853.
                                                                                                                                                                                                                                                                                  Length 13;
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                            Claim 1; SEQ ID NO 10370; 29pp + Sequence Listing; German.
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                                                                                                                                                                                                                                                       Sequence 13 BP; 5 A; 5 C; 1 G; 2 T; 0 U; 0 Other;
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Pred. No. 1.3e+03;
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Best Local Similarity 90.9%;
Matches 10; Conservative
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 methylation status
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data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                        Sequence 13 BP; 10 A; 3 C; 0 G; 0 T; 0 U; 0 Other;
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Matches 10; Conservative
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07-APR-2000; 2000DE-01019173.
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                                                                                                                         SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                      Oligonucleotide SEQ ID NO 86673 for detecting SNP TSC0021775.
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                                         ABC86656 standard; DNA; 13
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and merabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo int/pub/published_pct_sequences
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peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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13 TTTGTTTGGTTTY 1
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                                                                                                         Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 2422
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Matches
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scruttzi-899.rng

W0200177384-A2. Homo sapiens. 18-OCT-2001. 13 ABF42385; Query Match RESULT 2424 Best Loca Matches ABF42385 88333333333333388 g ò ö This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99889, ABF00010-ABF99889, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic. Gaps Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status. ö Oligonucleotide SEQ ID NO 132539 for detecting SNP TSC0033059. Claim 1; SEQ ID NO 125458; 29pp + Sequence Listing; German. 12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03; ive 0; Mismatches 1; Indels Sequence 13 BP; 7 A; 2 C; 0 G; 4 T; 0 U; 0 Other; Berlin K; ABF32542 standard; DNA; 13 BP. 06-APR-2001; 2001WO-IB000713. (first entry) Best Local Similarity 90.9 Matches 10; Conservative 943 ATTGGTTTAAT 953 Olek A, Piepenbrock C, 12 ATTGGATTAAT 2 (EPIG-) EPIGENOMICS AG WPI; 2001-657177/75 WO200177384-A2 Homo sapiens 21-FEB-2002 18-OCT-2001 ABF32542; Query Match

8 8

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretraeted genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic discorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99899, ABF00010-ABF99989, ABH0010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo int/pub/published_pct_sequences acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically precreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and range of diseases including immune system, gastrointestinal, respiratory. Central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99889, ABC0010-ABP9989, ABC0010-ABP9989 and ABL0010-ABB9989 and ABI0010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic. Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status. ; 0 Oligonucleotide SEQ ID NO 142382 for detecting SNP TSC0035690. Claim 1; SEQ ID NO 142382; 29pp + Sequence Listing; German. 12.9%; Score 9.4; DB 1; Length 13; larity 90.9%; Pred. No. 1.3e+03; Conservative 0; Mismatches 1; Indels Sequence 13 BP; 1 A; 4 C; 0 G; 8 T; 0 U; 0 Other; Seguence 13 BP; 4 A; 1 C; 8 G; 0 T; 0 U; 0 Other; Berlin K; ВР 06-APR-2001; 2001WO-IB000713. 07-APR-2000; 2000DE-01019173. ABF42385 standard; DNA; 13 21-FEB-2002 (first entry) 932 CCCTCCTCTC 942 Olek A, Piepenbrock C, m (EPIG-) EPIGENOMICS AG CCCTCCTCGTC WPI; 2001-657177/75 Local Similarity les 10; Conserv

This invention describes novel oligonuclectide primers or peptide nucleic

Claim 1; SEQ ID NO 132539; 29pp + Sequence Listing; German.

Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.

Berlin

Olek A, Piepenbrock C,

WPI; 2001-657177/75.

(EPIG-) EPIGENOMICS AG.

07-APR-2000; 2000DE-01019173

Query Match

Best Loca Matches

ABF43102;

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo.int/pub/published_pct_sequences
                                                                                                                                        SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                  Oligonucleotide SEQ ID NO 193268 for detecting SNP TSC0047550.
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                                                         (first entry)
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                                                         22-FEB-2002
                                                                                                                                                                                                                             Homo sapiens
                   ABF93271;
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                                                                                                                                                                                                                                                                                                                                                                                                             SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                                                                                                      Oligonucleotide SEQ ID NO 143099 for detecting SNP TSC0035891.
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                 Score 9.4; DB 1; Length 13;
Pred. No. 1.3e+03;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 9.4; DB 1; Length 13; Pred. No. 1.3e+03; 0; Mismatches 1; Indels
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ABF93271/c
ID ABF93271 standard; DNA; 13 BP.
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                 12.9%;
90.9%;
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90.9%;
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                                       Similarity
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                        Local br. 10;
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olek A,

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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
Oligonucleotide SEQ ID NO 220280 for detecting SNP TSC0053606.
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Gaps

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Page 1057
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABE99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic. 0; Gaps Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status. Oligonucleotide SEQ ID NO 195994 for detecting SNP TSC0048213. Claim 1; SEQ ID NO 220280; 29pp + Sequence Listing; German. Query Match 12.9%; Score 9.4; DB 1; Length 13; Best Local Similarity 90.9%; Pred. No. 1.3e+03; Matches 10; Conservative 0; Mismatches 1; Indels Sequence 13 BP; 4 A; 2 C; 0 G; 7 T; 0 U; 0 Other; Berlin BP. 06-APR-2001; 2001WO-IB000713. 07-APR-2000; 2000DE-01019173. 06-APR-2001; 2001WO-IB000713. 07-APR-2000; 2000DE-01019173 ABF95997 standard; DNA; 13 (first entry) 948 TITAATGIATC 958 Olek A, Piepenbrock C, TTTAATATATC 12 (EPIG-) EPIGENOMICS AG (EPIG-) EPIGENOMICS WPI; 2001-657177/75 WO200177384-A2 WO200177384-A2 Homo sapiens 22-FEB-2002 18-OCT-2001 18-OCT-2001 ABF95997; RESULT 2428 ò

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABE99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI32073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                     Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                                         German.
                                                                                                                      SEQ ID NO 195994; 29pp + Sequence Listing;
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.; 0 12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03; ive 0; Mismatches 1; Indels Sequence 13 BP; 0 A; 7 C; 2 G; 4 T; 0 U; 0 Other; Local Similarity 90.5 nes 10; Conservative Query Match Best Loca Matches

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Gaps

932 CCCTCCTTC 942 ||||||||||||||||2 2 CCCTCGTCTTC 12 셤 ò

Oligonucleotide SEQ ID NO 172077 for detecting SNP TSC0005766. ABF7208072
ID ABF72080 standard; DNA; 13 BP. AC ABF72080;
XX ABF72080;
XX 22-FEB-2002 (first entry)
XX 22-FEB-2002 (first entry)
XX SNP; single nuclectide polymorphic pertural nervous system; gastroint max with peptide nucleic acid; cytosine max with peptide nucleic acid; cytosine max with peptide nucleic acid; cytosine max with w0200177384-A2.
XX Homo sapiens.
XX PN W0200177384-A2.
XX SP 06-APR-2000; 2000DB-01019173.
XX PR 07-APR-2000; 2000DB-01019173.
XX SP 01-APR-2000; 2000DB-01019173.
XX WPI; 2001-657177/75.
XX Set of oligonucleotides, useful for Set of oligonucleotides, useful for Set of oligonucleotides, useful for Set of oligonucleotides including immunic cand cytosine methylation status:
XX Claim 1; SEQ ID NO 172077; 29pp for discontine methylation status:
XX Claim 1; SEQ ID NO 172077; 29pp for discontine methylation status:
XX Claim 1; SEQ ID NO 172077; 29pp for discontine methylation status:
XX Claim 1; SEQ ID NO 172077; 29pp for discontine methylation status:
XX Claim 1; SEQ ID NO 172077; 29pp for discontine methylation status:
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XX Claim 1; SEQ ID NO 172077; 29pp for discontine methylation status:
XX Claim 1; SEQ ID NO 172077; 29pp for discontine methylation status:
XX Claim 1;

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Berlin K;

Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.

Claim 1; SEQ ID NO 172077; 29pp + Sequence Listing; German.

This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The

Berlin K;

Olek A, Piepenbrock C,

WPI; 2001-657177/75

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oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF99889, ABH00010-ABH99989 and ABI00010-ABH8073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at frow poolity pub/published_pct_sequences
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                                                                                                                                                                                                                                                                                                          Score 9.4; DB 1; Length 13; Pred. No. 1.3e+03; 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                Sequence 13 BP; 9 A; 0 C; 3 G; 1 T; 0 U; 0 Other;
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Matches 10; Conservative
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                                                                                                                                                                                                                                                                                                                         Query Match
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                                                                                                                                                                                                                                     SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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designed to detect single-nucleotide polymorphisms and cytosine
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                                                                                                                                                                                                        Oligonucleotide SEQ ID NO 173482 for detecting SNP TSC0043214.
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76.9%;
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947 GTTTAATGTAT 957
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                              1 GTTTAACGTAT 11
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21-FEB-2002 (first entry)

Berlin K;

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically precreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABE09989, ABF00010-ABE99989, ABF00010-ABE99989, ABF00010-ABE99989, ABF00010-ABE99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at the printed specification, but ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                        Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                                                                                                                            Claim 1; SEQ ID NO 226938; 29pp + Sequence Listing; German.
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             07-APR-2000; 2000DE-01019173.
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                                                                          Olek A, Piepenbrock C,
                                              (EPIG-) EPIGENOMICS
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ABH27292/4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                           SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardicvascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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 Oligonucleotide SEQ ID NO 148524 for detecting SNP TSC0037485.
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Pred. No. 1.3e+03;
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Best Local Similarity 90.9%;
Matches 10; Conservative (
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                                                                                                 Homo sapiens
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                          This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretracted genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for ABH00010-ABH99989 and ABH00010-ABH99989, ABF00010-ABH99989, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at
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  SEQ ID NO 227269; 29pp + Sequence Listing; German.
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Pred. No. 1.3e+03;
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Best Local Similarity
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  Claim 1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                Length 13;
                                                                                                                          1; Indels
Sequence 13 BP; 3 A; 6 C; 0 G; 3 T; 0 U; 1 Other;
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                                                      Score 9.4; DB 1;
Pred. No. 1.3e+03;
0; Mismatches 1;
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76.9%;
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                                                             Query Match
Best Local Similarity 90.9%;
Matches 10; Conservative
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Best Local Similarity 76.9°
Matches 10; Conservative
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ABH28111 standard; DNA; 13 BP.

ABH28111

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schultzi-899.rng
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WO200177384-A2

06-APR-2001; 2001WO-IB000713.

07-APR-2000; 2000DE-01019173

(EPIG-) EPIGENOMICS

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomera are also used for detecting cell type differentiation. ABC00010-ABE09989, ABF00010-ABE99989, ABF00010-ABE99989 and ABI00010-ABI32073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at

0; Gaps Query Match 12.9%; Score 9.4; DB 1; Length 13; Best Local Similarity 90.9%; Pred. No. 1.3e+03; Matches 10; Conservative 0; Mismatches 1; Indels

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944 TTGGTTTAATG 954 | ||||||||| TAGGTTTAATG 12 N ઠ g

This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at

Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.

Berlin K;

Olek A, Piepenbrock C,

WPI; 2001-657177/75.

(EPIG-) EPIGENOMICS AG

06-APR-2001; 2001WO-IB000713. 07-APR-2000; 2000DE-01019173.

WO200177384-A2. Homo sapiens

18-OCT-2001.

Claim 1; SEQ ID NO 228088; 29pp + Sequence Listing; German.

RESULT 2439

Oligonucleotide SEQ ID NO 229254 for detecting SNP TSC0055935.

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Oligonucleotide SEQ ID NO 229253 for detecting SNP TSC0055935.

BP.

ABH29276 standard; DNA; 13

RESULT 2438

22-FEB-2002 (first entry)

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ABH29276;

Homo sapiens.

18-OCT-2001

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Oligonucleotide SEQ ID NO 228088 for detecting SNP TSC0055622.

(first entry)

22-FEB-2002

ABH28111;

Berlin K; Olek A, Piepenbrock C,

WPI; 2001-657177/75.

Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.

Claim 1; SEQ ID NO 229253; 29pp + Sequence Listing; German.

Sequence 13 BP; 4 A; 0 C; 3 G; 6 T; 0 U; 0 Other;

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0; Gaps

Query Match
12.9%; Score 9.4; DB 1; Length 13;
Best Local Similarity 90.9%; Pred. No. 1.3e+03;
Matches 10; Conservative 0; Mismatches 1; Indels

943 ATTGGTTTAAT 953

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11 Argedriraar 1

Sequence 13 BP; 6 A; 3 C; 0 G; 4 T; 0 U; 0 Other;

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Berlin K;

Piepenbrock C,

olek A,

designed to detect methylation status.

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Gaps

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The
                                                                                                                                                                                                                       Claim 1; SEQ ID NO 207761; 29pp + Sequence Listing; German.
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Berlin K;

Piepenbrock C,

olek A,

WPI; 2001-657177/75

(EPIG-) EPIGENOMICS AG

06-APR-2001; 2001WO-IB000713 07-APR-2000; 2000DE-01019173

WO200177384-A2 Homo sapiens

18-OCT-2001.

12.9%; Score 9.4; DB 1; Length 13;

Query Match

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABF92073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but two botained in electronic format from WIPO at
oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The coligoners are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                               12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03;
                                                                                                                                                                                                                                                                                                           1; Indels
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                                                                                                                                                                                                                                                                                                             0; Mismatches
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Matches 10, Conservative
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                                                                                                                                                                                                                                        This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                   Claim 1; SEQ ID NO 229254; 29pp + Sequence Listing; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ch 12.9%; Score 9.4; DB 1; Length 13; 1 Similarity 90.9%; Pred. No. 1.38+03; 10; Conservative 0; Mismatches 1; Indels
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BP.

ABH07784 standard; DNA; 13

944 TIGGITIPATG 954

Query Match Best Local Similarity

Matches

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12 raggrirants

(first entry)

22-FEB-2002

ABH07784;

was obtained in electronic format from Wl ftp.wipo.int/pub/published_pct_sequences

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                                                                                                                                       SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
        Gaps
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                                                                                                                          Oligonucleotide SEQ ID NO 158090 for detecting SNP TSC0039821.
Pred. No. 1.3e+03;
; Mismatches 1; Indels
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90.9%;
                                                                             ABF58093 standard; DNA; 13
                                                                                                            (first entry)
Best Local Similarity 90.9
Matches 10; Conservative
                       905 TCATTTTCTT
                                     12 rCATATICITI
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                                                                                                                                                                          Homo sapiens.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and oytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABH99989 and ABI00010-ABI82073 invepresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at Etp.wipo.int/pub/published_pct_sequences

Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine methylation status.

ĸ Berlin

Olek A, Piepenbrock C, (EPIG-) EPIGENOMICS AG

WPI; 2001-657177/75.

Claim 1; SEQ ID NO 158090; 29pp + Sequence Listing; German.

Sequence 13 BP; 8 A; 2 C; 1 G; 1 T; 0 U; 1 Other;

.; 0 12.9%; Score 9.4; DB 1; Length 13; larity 76.9%; Pred. No. 1.3e+03; Conservative 1; Mismatches 2; Indels Best Local Similarity Matches 10; Conserv Query Match à

ABF85688 standard; DNA; 13 RESULT 2443 ABF85688

ABF85688

BP.

Oligonucleotide SEQ ID NO 185685 for detecting SNP TSC0045759. (first 22-FEB-2002

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Homo sapiens.

WO200177384-A2.

18-OCT-2001

06-APR-2001; 2001WO-IB000713

07-APR-2000; 2000DE-01019173

(EPIG-) EPIGENOMICS

Berlin K; Piepenbrock C, olek A,

WPI; 2001-657177/75

set or oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.

Claim 1; SEQ ID NO 185685; 29pp + Sequence Listing; German.

acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomers located in chemically pretreated genomic DNA. The oligomers are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gasteminestinal, respiratory, cingomers are also used for detecting cell type differentiation. ABCC0010-ABC99889, ABC0010-ABC99889, ABC0010-ABC99889 ABC0010-ABC99889 and ABI0010-ABC80010-ABC6010 ABC6010 This invention describes novel oligonuclectide primers or peptide nucleic

Sequence 13 BP; 2 A; 0 C; 4 G; 6 T; 0 U; 1 Other;

Gaps . 0 12.9%; Score 9.4; DB 1; Length 13; 76.9%; Pred. No. 1.3e+03; ative 1; Mismatches 2; Indels Query Match 12.9 Best Local Similarity 76.9 Matches 10; Conservative

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945 TGGTTTAATGTAT 957 1 recrirationax 13 D à

657/c ABH11657 standard; DNA; 13

ABH11657;

22-FEB-2002 (first entry)

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Gaps

Oligonucleotide SEQ ID NO 211634 for detecting SNP TSC0051609.

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Homo sapiens.

WO200177384-A2.

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07-APR-2000;

olek A,

18-OCT-2001

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                                                                                                               This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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  designed to detect single-nucleotide polymorphisms and cytosine
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                                                                     Claim 1; SEQ ID NO 212324; 29pp + Sequence Listing; German.
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Pred. No. 1.3e+03;
1; Mismatches 2; Indels
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76.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                     This invention describes novel oligonucleotide primers or peptide nucleic
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                                                                                                                                                                                                                                                                                    Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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Query Match

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
represent the oligomers described in the invention. NOTE: The sequence data for this parent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pot_sequences
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12.9%; Score 9.4; DB 1; Length 13;
Best Local Similarity 90.9%; Pred. No. 1.3e+03;
Matches 10; Conservative 0; Mismatches 1; Indels
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12.9%; Score 9.4; DB 1; Length 13;
Best Local Similarity 90.9%; Pred. No. 1:38+03;
Matches 10; Conservative 0; Mismatches 1; Indels
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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Oligonucleotide SEQ ID NO 241532 for detecting SNP TSC0058904

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                      Piepenbrock C,
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                              (EPIG-) EPIGENOMICS AG.
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                                                                                      olek A,
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Berlin K;
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                                                                                                                        Homo sapiens
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Oligonucleotide SEQ ID NO 253873 for detecting SNP TSC0061899.
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Sequence 13 BP; 1 A; 0 C; 4 G; 8 T; 0 U; 0 Other;

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence was obtained in electronic format from MIPO at the printed specification, but the wipo.int/pub/published_pct_sequences
This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99889, ABF00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at the printed specification, but fire wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Best Local Similarity 76.9
Matches 10; Conservative
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ABH58822
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Length 13;
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Score 9.4; DB 1;
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ftp.wipo.int/pub/published_pct_sequences
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12.9%;
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Best Local Similarity 90.9
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Query Match 12.9
Best Local Similarity 90.9
Matches 10; Conservative
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                                                                      913 TTTGGTCTTTG 923
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RESULT 2454 ABC42528

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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic formmat from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                           Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                                                                                                                        (EPIG-) EPIGENOMICS AG
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                                                                                                                                                                                                         SNP, single nucleotide polymorphism, human; diagnosis; PNA, cancer, CNS, peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                           Oligonucleotide SEQ ID NO 42545 for detecting SNP TSC0012678.
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ABC42528 standard; DNA; 13
                                                                                                     (first entry)
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Best Local Similarity 90.9
Matches 10; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           methylation status.
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                              Gaps
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 12.9%; Score 9.4; DB 1; Length 13; 76.9%; Pred. No. 1.3e+03; Attive 1; Mismatches 2; Indels
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                                                                                                                                                    ABC99599 standard; DNA; 13 BP.
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Query Match
Best Local Similarity 76.9°
Matches 10, Conservative
                                                                          13 TTTGTCGTTTATY 1
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                                                                                                                          This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretraeted genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABH99989 and ABI00010-ABH82073 tarpresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but they wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                              Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                               Claim 1; SEQ ID NO 99616; 29pp + Sequence Listing; German.
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Best Local Similarity 90.9
Matches 10; Conservative
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central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABH99989 and ABI00010-ABI32073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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ftp.wipo.int/pub/published_pct_sequences
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ilarity 76.9%;
Conservative
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Best Local Similarity
Matches 10; Conserv
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Oligonucleotide SEQ ID NO 2892 for detecting SNP TSC0001129.

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                                                                                                                                                                                                                                   SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Pred. No. 1.3e+03;
0; Mismatches 1; Indels
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90.9%;
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GGTTTAATGTATC 958
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                    13 GGTTTAATGGGTY
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 trepresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                         SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; SEQ ID NO 2892; 29pp + Sequence Listing; German.
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                                                                                                                                                                                                                                                                                                                   Berlin K;
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12 TTTGCGTTTTA
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Best Local Similarity
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                                                                                                                Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                               Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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12.9%; Score 9.4; DB 1; Length 13;
Best Local Similarity 76.9%; Pred. No. 1.3e+03;
Matches 10; Conservative 1; Mismatches 2; Indels
                                                                                                                                                                                                                                                           Claim 1; SEQ ID NO 30479; 29pp + Sequence Listing; German.
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06-APR-2001; 2001WO-IB000713.
                                     07-APR-2000; 2000DE-01019173.
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                                                                      (EPIG-) EPIGENOMICS AG
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99999, ABF00010-ABF99999, ABH00010-ABH99999 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIFO at ftp.wipo.int/pub/published_pct_sequences
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Claim 1; SEQ ID NO 105982; 29pp + Sequence Listing; German.
                                                                                                                                                                                                                                                                                                                                                                                            12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03; ive 0; Mismatches 1; Indels
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947 GITTAATGTAT 957
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                                                                                                                                                                                                                                                                                                                                                                                                                             SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                 12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03; ive 0; Mismatches 1; Indels
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                                                 Sequence 13 BP; 7 A; 0 C; 4 G; 2 T; 0 U; 0 Other;
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was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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Best Local Similarity
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ABC58137/c
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99899, ABF00010-ABF99989, ABF00010-ABF99989, ABF00010-ABF99989, abmoost the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPD at
                                                                                                                                                                                                                                                                                                         SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                           Oligonucleotide SEQ ID NO 9258 for detecting SNP TSC0002455.
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Pred. No. 1.3e+03;
0; Mismatches 1; Indels
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                                ABC09267 standard; DNA; 13 BP.
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les 10; Conservative
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                                                                                                     ABC09267;
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central nervous system; gastrointestinal; respiratory; immune; metabolic.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99999, ABF00010-ABF99999, ABH00010-ABH99999 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                                                       Claim 1; SEQ ID NO 85843; 29pp + Sequence Listing; German.
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Best Local Similarity 90.5
Matches 10; Conservative
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and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABR0010-ABC9989, ABR0010-ABS9989, ABR0010-ABS9989 and ABI0010-ABI82073 represent the oligomers described in the invention. NoTE: The sequence data for this patent did not form part of the printed specification, but ftp.wipo.int/pub/published_pct_sequences
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Pred. No. 1.3e+03;
0; Mismatches 1; Indels
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90.9%;
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Sequence 13 BP; 2 A; 0 C; 2 G; 8 T; 0 U; 1 Other;

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Score 9.4; DB 1; Length 13; Pred, No. 1.3e+03; 1; Mismatches 2; Indels
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Best Local Similarity 76.9
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ABF15751/c
ID ABF1575
XX
                                                                                                                                                                                                                                                                                                            RESULT 2470
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ID ABC40251
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC09989, ABF00100-ABF9989, ABF00100-ABF9989, ABF00100-ABF9989, ABF00100-ABF9989, and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at
                                                                                                  SNP; single nuclectide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                   Oligonucleotide SEQ ID NO 115748 for detecting SNP TSC0029020.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 1; SEQ ID NO 115748; 29pp + Sequence Listing; German.
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                                                                                                                                                                                                                                                                                       06-APR-2001; 2001WO-IB000713.
                                                                                                                                                                                                                                                                                                                            07-APR-2000; 2000DE-01019173.
                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 12.9
Best Local Similarity 90.9
Matches 10; Conservative
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                                21-FEB-2002
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC09989, ABF00010-ABF99989, ABH0010-ABF99989, and ABI00110-ABF8073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic formmat from WIPO at

Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.

Berlin K;

Olek A, Piepenbrock C,

(EPIG-) EPIGENOMICS

WPI; 2001-657177/75

06-APR-2001; 2001WO-IB000713 07-APR-2000; 2000DE-01019173

18-OCT-2001

Claim 1; SEQ ID NO 42131; 29pp + Sequence Listing; German.

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12.9%; Score 9.4; DB 1; Length 13; 76.9%; Pred. No. 1.3e+03; tive 1; Mismatches 2; Indels

946 GGTTTAATGTATC 958

Query Match
Best Local Similarity 76.9
Matches 10, Conservative

GGTTTGATGTTTY 13

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ABC66733 standard; DNA; 13

21-FEB-2002 (first entry)

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ABC66733;

Sequence 13 BP; 1 A; 0 C; 4 G; 7 T; 0 U; 1 Other;

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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Oligonucleotide SEQ ID NO 66750 for detecting SNP TSC0017501.
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RESULT 2473
ABC66733/C
XX
AC ABC66733/C
AC ABC66733/C
DT 21-FEB-7
XX
DT 21-FEB-7
XX
CONTROL SUP, SIP
XX
CONTROL SUP
XX
CONTROL
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Berlin K; Piepenbrock C, WPI; 2001-657177/75 Olek A,

07-APR-2000; 2000DE-01019173

(EPIG-) EPIGENOMICS AG

ABC42114 standard; DNA; 13 BP. ABC4211 BXSXXXXXXXXXXXXXX

ABC42114;

Oligonucleotide SEQ ID NO 42131 for detecting SNP TSC0012592.

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Homo sapiens

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RESULT 2472

21-FEB-2002 (first entry)

WO200177384-A2

-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences

88888888

Gaps

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ch 12.9%; Score 9.4; DB 1; Length 13; 1. Similarity 90.9%; Pred. No. 1.38+03; 10; Conservative 0; Mismatches 1; Indels

Query Match Best Local S: Matches 10

934 CTCCTCTTCAT 944

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crecregrear 1

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Sequence 13 BP; 4 A; 1 C; 5 G; 2 T; 0 U; 1 Other;

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; SEQ ID NO 119571; 29pp + Sequence Listing; German.
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                                                                                                                                                                                                                                                                                                                   acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABR99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                  This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP)
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Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                                                                                 SEQ ID NO 66750; 29pp + Sequence Listing; German.
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
Oligonucleotide SEQ ID NO 119571 for detecting SNP TSC0029845.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABE99989, ABF00010-ABE99989, ABF00010-ABE99989 and ABI00010-ABE82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WFPO at fitted specification, but ftp.wipo.int/pub/published_pct_sequences ö Gaps . 0 Score 9.4; DB 1; Length 13; Pred. No. 1.3e+03; 0; Mismatches 1; Indels Sequence 13 BP; 3 A; 0 C; 1 G; 9 T; 0 U; 0 Other; 12.9%; 10; Conservative Query Match Best Local Similarity Matches

947 GTTTAATGTAT 957

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Homo sapiens

WO200177384-A2.

18-OCT-2001.

06-APR-2001; 2001WO-IB000713.

07-APR-2000; 2000DE-01019173

(EPIG-) EPIGENOMICS AG

Berlin Piepenbrock C, olek A,

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WPI; 2001-657177/75

Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine methylation status.

Claim 1; SEQ ID NO 120579; 29pp + Sequence Listing; German.

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ABF20582 standard; DNA; 13 BP.

ABF20582;

(first entry) 21-FEB-2002 Oligonucleotide SEQ ID NO 120579 for detecting SNP TSC0030084.

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                      Oligonucleotide SEQ ID NO 125459 for detecting SNP TSC0031370.
                                                                                                                                                                                                                                     Berlin K;
                                      ABF25462 standard; DNA; 13 BP.
                                                                                                                                                                                     06-APR-2001; 2001WO-IB000713.
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                                                                     (first entry)
                                                                                                                                                                                                                                     Olek A, Piepenbrock C,
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                                                                                                                                                     WO200177384-A2
                                                                                                                                       Homo sapiens.
                                                                      21-FEB-2002
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                                                       ABF25462;
                       RESULT 2476
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Set of oligonuclectides, useful for diagnosis and cell typing, idesigned to detect single-nuclectide polymorphisms and cytosine methylation status.

Berlin K;

Olek A, Piepenbrock C, (EPIG-) EPIGENOMICS

WPI; 2001-657177/75.

06-APR-2001; 2001WO-IB000713. 07-APR-2000; 2000DE-01019173.

WO200177384-A2

18-OCT-2001

Homo sapiens,

Claim 1; SEQ ID NO 128732; 29pp + Sequence Listing; German.

Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status. Claim 1; SEQ ID NO 125459; 29pp + Sequence Listing; German. ftp.wipo.int/pub/published_pct_sequences

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at Sequence 13 BP; 3 A; 0 C; 3 G; 7 T; 0 U; 0 Other;

. 0 12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03; ive 0; Mismatches 1; Indels Query Match Best Local Similarity 90.9 Matches 10; Conservative

ABF32538

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Gaps

943 ATTGGTTTAAT 953 ATTGGGTTAAT 12

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ABF28735 standard; DNA; 13 ABF28735 2477 RESULT 24'
ABF28735/A
ID ABF28
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AC ABF2
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DT 21-FI
XX
DE Olige

BP.

Oligonucleotide SEQ ID NO 128732 for detecting SNP TSC0032227.

(first entry)

21-FEB-2002

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                                                                                                                                                                                                                                                                                                                                                                                                     This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from MIPO at
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hes 10; Conservative
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic. Oligonucleotide SEQ ID NO 132535 for detecting SNP TSC0033059. (first entry) Homo sapiens 21-FEB-2002

06-APR-2001; 2001WO-IB000713 WO200177384-A2 18-OCT-2001.

07-APR-2000; 2000DE-01019173.

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF9989, ABF00010-ABF9989, ABF00010-ABF9989, and ABI0010-ABF82073 represent the oligomers described in the invention. NoTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from NIPO at
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designed to detect single-nucleotide polymorphisms and cytosine
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Pred. No. 1.3e+03;
0; Mismatches 1; Indels
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                                                                                                                             Berlin K;
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                                                              (EPIG-) EPIGENOMICS AG
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Claim 1; SEQ ID NO 132536; 29pp + Sequence Listing; German.

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                This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC09989, ABF00010-ABF99899, ABF00010-ABF99899 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo.int/pub/published_pct_sequence
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic. Oligonucleotide SEQ ID NO 193663 for detecting SNP TSC0047644. BP. ABF93666 standard; DNA; 13 (first entry) 22-FEB-2002 ABF93666; 2483 ABF93666 ò 음

ABH18280 standard; ABH18280;

ABH18280

BP.

DNA; 13

22-FEB-2002 (first entry)

Oligonucleotide SEQ ID NO 218257 for detecting SNP TSC0053052.

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Homo sapiens,

WO200177384-A2

18-OCT-2001

06-APR-2001; 2001WO-IB000713.

07-APR-2000; 2000DE-01019173.

(EPIG-) EPIGENOMICS AG.

Berlin K; Olek A, Piepenbrock C,

WPI; 2001-657177/75.

Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.

Claim 1; SEQ ID NO 218257; 29pp + Sequence Listing; German.

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABE99989, ABF00010-ABE99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at

Sequence 13 BP; 3 A; 0 C; 3 G; 7 T; 0 U; 0 Other;

12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03; ive 0; Mismatches 1; Indels 10; Conservative Local Similarity Query Match Matches

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, contral nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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ftp.wipo.int/pub/published_pct_sequences
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated gendmic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABE9989, ABF00010-ABE9989, ABF00010-ABE9989, ABF00010-ABE9989 and ABI00010-ABI82073 trepresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00100-ABC099889, ABC00100-ABE99889 and ABI00010-ABE99889 and ABI00010-ABE99899 and ABI00010-ABE9073 arepresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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12.9%; Score 9.4; DB 1; Length 13;
Best Local Similarity 76.9%; Pred. No. 1.38+03;
Matches 10; Conservative 1; Mismatches 2; Indels
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12.9%; Score 9.4; DB 1; 90.9%; Pred. No. 1.3e+03;

Query Match Best Local Similarity Page 1082

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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99999, ABF00010-ABF99999, ABH00010-ABH99999 and ABI00010-ABF3073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo.int/pub/published_pct_sequences
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designed to detect single-nucleotide polymorphisms and cytosine
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                                                                                                                                SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                 Oligonucleotide SEQ ID NO 227328 for detecting SNP TSC0004949.
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Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine

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                                                       This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and oytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF001010-ABF9989, ABF001010-ABF9989, ABF001010-ABF9989 and ABI00110-ABF82073 represent the oligomers described in the invention. NOTE: The sequence was obtained in electronic format from WIPO at the printed specification, but the wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                Claim 1; SEQ ID NO 203266; 29pp + Sequence Listing; German.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABP00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI22073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from NIPO at
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data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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Best Local Similarity 90.9%; Pred. No. 1.3e+03;
Matches 10; Conservative 0; Mismatches 1; Indels
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Pred. No. 1.3e+03;
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                                                                                                                                Seguence 13 BP; 4 A; 0 C; 1 G; 7 T; 0 U; 1 Other;
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90.9%;
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Best Local Similarity 90.9
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                                                                                                                                                          SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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designed to detect single-nucleotide polymorphisms and cytosine
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RESULT 2493
ABH07785/c
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo.int/pub/published_pct_sequences
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peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Pred. No. 1.3e+03;
0; Mismatches 1; Indels
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90.9%;
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Best Local Similarity 90.9
Matches 10, Conservative
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Berlin K; Olek A, Piepenbrock C, EPIGENOMICS AG WPI; 2001-657177/75. (EPIG-)

set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.

Claim 1; SEQ ID NO 236753; 29pp + Sequence Listing; German.

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at

Sequence 13 BP; 10 A; 0 C; 2 G; 1 T; 0 U; 0 Other;

0; Gaps 12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03; rive 0; Mismatches 1; Indels Local Similarity 90.9 Query Match

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ABF61733 standard; DNA; 13 BP. ABF61733; RESULT 2496 ABF61733/
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XXY SIP,
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XXX Gent
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YX

(first entry) 22-FEB-2002 Oligonucleotide SEQ ID NO 161730 for detecting SNP TSC0040712.

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Homo sapiens.

WO200177384-A2

18-OCT-2001,

06-APR-2001; 2001WO-IB000713.

07-APR-2000; 2000DE-01019173.

(EPIG-) EPIGENOMICS AG.

Berlin Olek A, Piepenbrock C, WPI; 2001-657177/75.

Set of oligonuclectides, useful for diagnosis and cell typing, i designed to detect single-nuclectide polymorphisms and cytosine designed to detect methylation status.

Claim 1; SEQ ID NO 161730; 29pp + Sequence Listing; German.

This invention describes novel oligonucleotide primers or peptide nucleic

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acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99889, ABC0010-ABC99889, ABC0010-ABC99889 and ABI0010-ABC80010 crepsent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPD at fitp.wipo.int/pub/published_pct_sequences
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Sequence 13 BP; 7 A; 3 C; 1 G; 1 T; 0 U; 1 Other;

Gaps .; 0 Length 13; Score 9.4; DB 1; Length 13 Pred. No. 1.3e+03; 1; Mismatches 2; Indels 12.9%; 76.9%; Query Match
Best Local Similarity 76.9
Matches 10, Conservative

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1948 TITAATGTATCGC 960 ||||| ||| |||: TTTATTGTGTCGY 1 13

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RESULT 24 ABF62775/

ABF62775 standard; DNA; 13

BP. ABF62775; Oligonucleotide SEQ ID NO 162772 for detecting SNP TSC0040932.

(first entry)

22-FEB-2002

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SND; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Homo sapiens.

WO200177384-A2

18-OCT-2001

06-APR-2001; 2001WO-IB000713.

07-APR-2000; 2000DE-01019173

(EPIG-) EPIGENOMICS AG.

Berlin K; Olek A, Piepenbrock C, WPI; 2001-657177/75.

Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.

Claim 1; SEQ ID NO 162772; 29pp + Sequence Listing; German.

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99899, ABH00010-ABH99899 and ABT00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from MIPO at ftp.wipo.int/pub/published_pct_sequences

Sequence 13 BP; 4 A; 3 C; 0 G; 6 T; 0 U; 0 Other;

Oligonucleotide SEQ ID NO 163721 for detecting SNP TSC0041134.

(first entry)

22-FEB-2002

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                                                                                                                                                                                                                                                                                                                                     SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Score 9.4; DB 1; Length 13;
Pred. No. 1.3e+03;
0; Mismatches 1; Indels
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                                     10; Conservative
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Query Match
Best Local Similarity
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 the represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                    SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Matches 10; Conservative
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Homo sapiens,

ABF63724 standard; DNA; 13 BP.

RESULT 2499 ABF63724 ID ABF63724

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                            designed to detect methylation status.
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WO200177384-A2
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF9989, ABH0010-ABH9989 and ABI00010-ABI82073 trepresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from MIPO at
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Set of oligonuclectides, useful for diagnosis and cell typing, is designed to detect single-nuclectide polymorphisms and cytosine methylation status.
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                                                                                                                                                       Claim 1; SEQ ID NO 247681; 29pp + Sequence Listing; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12.9%; Score 9.4; DB 1; Length 13; 90.9%; Pred. No. 1.3e+03; ive 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 13 BP; 4 A; 0 C; 3 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             was obtained in electronic format from W.
ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Berlin K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ABH60732 standard; DNA; 13 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (EPIG-) EPIGENOMICS AG
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Best Local Similarity
Matches 10; Conserv
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The present invention describes a recombinant adeno-associated virus (AAV) vectored ribozyme composition (I). (I) comprises: (a) at least a first ribozyme that specifically cleaves an manA encoding a protein, polypeptide, or peptide selected from the group of rod opsin, iNOS, RDS/peripherin, VEGFR1, VEGFR2, adenosine A-2B receptor, IGF-1, integrin alpha 3, integrin alpha 5, not integrin alpha 7, integrin alpha 3, integrin alpha 5, or integrin alpha V, (b) a vector comprising a polymucleotide encoding the ribozyme, where the polymucleotide operably positioned downstream of at least a first promoter that directs expression of the polymucleotide in a selected mammalian cell transformed with the vector; (c) a viral particle comprising the ribozyme or the polymucleotide; (d) an AAV vector comprising the ribozyme or the polymucleotide; or (e) a host cell comprising the ribozyme or the polymucleotide, also described is a method for decreasing the amount of mRNA encoding a selected polypeptide in a composition described amammalian eye, comprising to the eye the composition described amammalian eye, comprising to the eye the composition described and man evel (I) has ophthalmological activity, and can
                                                                                                                                                                                                                                                                                                                                                                                                                    ö
oligomers are also used for detecting cell type differentiation. ABC000010-ABC9989, ABF00010-ABF9989, ABH00010-ABF9989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at fixed the printed specification, but ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hairpin ribozyme; hammerhead ribozyme; ribozyme; retinal disease; target; ophthalmological; gene therapy; eye; retinal dysfunction; AAV; diabetic retinopathy; macular degeneration; autosomal dominant retinitis; blood-retinal barrier dysfunction; adeno-associated virus; blindness; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A recombinant adeno-associated virus-vectored ribozyme composition, useful for treating a disease or dysfunction of the mammalian eye e.g. retinal disease, e.g. diabetic retinopathy or age-related macular
                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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                                                                                                                                                                                                                                                                                                                                      Score 9.4; DB 1; Length 13;
Pred. No. 1.3e+03;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                           Sequence 13 BP; 2 A; 0 C; 3 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   IGF1 Rz1 ribozyme target sequence SEQ ID NO:88.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-MAY-2002; 2002WO-US013679.
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                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 90.9
Matches 10, Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO200288320-A2.
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be used in gene therapy. (I) can be used for treating a disease or dysfunction of the mammalian eye, such as a retinal disease or retinal dysfunction, (diabetic) retinopathy, or (age-related) macular degeneration. (I) is also useful for manifacturing a medicament for treating the disease mentioned above, including autosomal dominant retinitis or a blood-retinal barrier dysfunction. (I) can also be useful for treating, decreasing the severity, or ameliorating the symptoms of a pathological condition, e.g. atrophic or pigmented lesions of the eye, blindness, a reduction in central or peripheral vision, or a reduction in total vision. ABZ72763 to ABZ72953 represent sequences used in the exemplification of the present invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Nucleic acid molecule, Hepatitis C virus, HCV; Hepatitis B virus, HBV; RNA stability, RNA expression, RNA synthesis, antieense; enzymatic nucleic acid, hammerhead ribozyme; DNAzyme; inozyme; zinzyme; amberzyme; G-cleaver ribozyme; decoy molecule; aptamer; HBV reverse transcriptase; Enhancer I region, viral replication; degenerative, disease state; HBV infection; HCV infection; cirrhosis; liver failure; hepatocellular carcinoma; hepatotropic; cytostatic; virucide, antiinflammatory; substrate; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HBV enzymatic nucleic acid substrate sequence #185.
                                                                                                                                                                                                                                                                                                                                                                       Sequence 13 BP; 1 A; 4 C; 2 G; 0 T; 6 U; 0 Other;
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08-UUN-2001, 2001US-00877478.
08-UUN-2001, 2001US-0296876P.
24-OCT-2001, 2001US-0337659P.
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| S UUCGUCUUUGC 12
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Roberts E;
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MACEJAK D.
MCSWIGGEN J.
MORRISSEY D.
PAVCO P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
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DRAPER K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ROBERTS E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO200281494-A1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
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(BLAT/)
(MACE/)
(MCSW/)
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(PAVC/)
(LEEP/)
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The present invention relates to nucleic acid molecules which modulate the synthesis, expression and/or stability of Hepatitis C virus (HCV) or Hepatitis B virus (HBV) RNA. The nucleic acid molecules include antisense and enzymatic nucleic acids such as hammerhead ribozymes, DNAzymes, or are nucleic acid decoy molecules and aptamers that bind to HBV reverse transcriptase that bind to HBV reverse transcriptase primer sequences, as well consolidation and aptamers that bind to HBV reverse transcriptase primer sequences, as well as oligonucleotides that specifically bind the Enhancer I region of HBV DNA. The nucleic acids may be used to modulate the expression of HBV C genes and HBV viral replication. Also disclosed is a method for screening compounds and/or potential therapies directed against HBV, and compounds that modulate the expression and/or replication of HCV. The compounds and compounds consolidation are useful for the treatment of degenerative and disease states related to HBV and HCV infection, replication and gene expression such as cirrhosis, liver failure, and hepatocellular carcinoma. The present sequence represents a substrate for one of the HBV enzymatic nucleic acid sequence represents a substrate for one of the HBV enzymatic nucleic acid sequences disclosed in the present invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Probes were prepared on the basis of two regions of the known sequence (Ogrydziak et al. J.Gen.Microbiol.(1982) 128,1255-1234) of the first 25 amino acids of mature extracellular alkaline protease. This probe is based on Region II, beginning at amino acid #18. It was hybridised with
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 9.4; DB 1; Length 13; Pred. No. 1.3e+03; 5; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Probe 186 to Yarrowia lipolytica XPR2 gene. (3'-5').
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 13 BP; 0 A; 5 C; 3 G; 0 T; 5 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  extracellular alkaline protease; XPR2; ss.
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                                                        Example 1; Page 221; 387pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example; Fig 2; 45pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            12.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            86EP-00307839.
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86US-00841121.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAN71348 standard; DNA; 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ouery Match
Best Local Similarity 45.5.
The state of the 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (PFIZ ) PFIZER INC. (USHU-) US HUIRUI CO LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            917 GICTITGCCTT 927
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Davidow LS, Franke AE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3 GUCUGUGCCUU 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (revised)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1987-124409/18.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            10-OCT-1986;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            18-OCT-1985;
18-MAR-1986;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              25-MAR-2003
25-APR-1991
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EP220864-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic.
infection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAN71348;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 2505
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AAV49008-236 represent antisense oligonucleotides directed against the rb gene. Of these, only oligonucleotides AAV49008-52 resulted in effective downrequlation of negative growth control by rb, while oligonucleotides AAV49052-26 had little effect. The oligonucleotides examplify the invention. The specification describes oligonucleotides that contain 8-30 nucleotides, which contain at most 8 nucleotides that contain 8-30 nucleotides, which contain at most 8 nucleotides that contain 8-30 nucleotides, which contain at most 8 nucleotides cach ach form three hydrogen bonds to cytosine, do not contain two sequences of three consecutive nucleotides ach able to form three H-bonds to three consecutive cytosines, and the ratio between three H-bonds to three consecutive cytosines, and the ratio between three H-bonds to form two H-bonds each [2R) or three such bonds (3R) is given by 2R/3R = 0.33-0.72. The oligonucleotides are used to modilate expression of genes, particularly the genes for p53, ErB-2, junB, junD, TGF-beta 1 or beta 2 to control proliferation of primary cell cultures and/or keratinocytes). The oligonucleotides can also be used to analyse therapeutically, e.g. in cases of cancer or (targeting TGF) for stimulating the immune system
                                                                                                                                                                                        ö
three overlapping plasmids recovered from the XPR2 transformant 'lipolytica Arca 20781 to confirm that the XPR2 gene had been cloned. See also AAN70213-NN0218, AAN71339, AAN71340, AAN71343-7. (Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Preparation of antisense oligo:nucleotide(s) which lack long runs of consecutive guanosine or inosine - and have specific ratio of residues able to form two or three hydrogen bonds, have greater activity and reduced toxicity, used therapeutically or to modulate growth of cells in
                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              rb gene; antisense oligonucleotide; modulate; gene expression; ss.
                                                                                                                                                                                      .;
0
                                                                                                                                                Length 14;
                                                                                                                                          Query Match 12.9%; Score 9.4; DB 1; Length 14 Best Local Similarity 71.4%; Pred. No. 1.4e+03; Matches 10; Conservative 1; Mismatches 3; Indels
                                                                                                    Sequence 14 BP; 0 A; 2 C; 3 G; 7 T; 0 U; 2 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (BIOG-) BIOGNOSTIK GES BIOMOLEKULARE DIAGNOSTIK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    rb gene antisense oligonucleotide rb-N-17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 7; Fig 9a; 286pp; English.
                                                                                                                                                                                                                                                                                                                                                                              AAV49069 standard; DNA; 14 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Brysch W;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  97EP-00101531
                                                                                                                                                                                                                               910 TICTITGGICTITG 923
                                                                                                                                                                                                                                                                     1 rrcrrycgngrrrg 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                             15-OCT-1998 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1998-400910/35.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Schlingensiepen K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  31-JAN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          05-AUG-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                     AAV49069;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 culture.
                                                                                                                                                                                                                                                                                                                                      RESULT 2506
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                                                                                                                                                                                                                                                                                                                                                                                    Triplex formation; DNA detection; triple helix; identification; bacteria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present sequence represents a polynucleotide that is able to form a triple helix with a double stranded sequence. Cytosine bases in the present can be replaced with 5-methylcytosine for increased triplex stability. The present sequence is used in the assay of the invention, where it can be part of the anchor DNA or reporter DNA sequence. The assay comprises adding a sample containing double-stranded DNA test sequences to an aqueous medium containing at least one complex of anchor DNA, attached to a solid support, and reporter DNA, where either a part of the anchor DNA or reporter DNA is designed to form a triple-strand
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              structure with part of the test sequence. Trippex formation results in displacement of the reporter DNA which is detected as an indication of the presence of the DNA test sequence. The method is used to detect DNA sequences, particularly for identification of bacteria (by detecting genes for ribosomal RNA) in clinical samples, but also detection of oncogenes and Hepatitis B virus
                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Assay of genetic sequences based on triplex formation from double stranded analyte - and hybrid of anchor and reporter sequences, with reporter released if triplex formation occurs, used e.g. to identify
                                                                                                                                                                                                                                                                                                                                                 Triple helix third strand of Prealbumin gene nucleotides 250-263.
                                                                         ö
                                Score 9.4; DB 1; Length 14; Pred. No. 1.4e+03; 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12.9%; Score 9.4; DB 1; Length 14; 90.9%; Pred. No. 1.4e+03; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 14 BP; 0 A; 6 C; 0 G; 8 T; 0 U; 0 Other;
Sequence 14 BP; 4 A; 2 C; 0 G; 8 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Col 17-18; 168pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (PROF-) PROFILE DIAGNOSTIC SCI INC.
                                    12.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        93US-00173489.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             92US-00968436.
                                                                                                                                                                                                                                      AAX14711 standard; DNA; 14
                                                                                                                                                                                                                                                                                                              24-MAR-1999 (first entry)
                                    Query Match 12.9
Best Local Similarity 90.9
Matches 10; Conservative
                                                                                                            905 TCATTTTCTTT 915
                                                                                                                                               2 rcaarrrcrrr 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hepburn AG, Wang C;
                                                                                                                                                                                                                                                                                                                                                                                                       oncogene; virus; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1999-130384/11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        22-DEC-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             29-OCT-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US5861244-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19-JAN-1999.
                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                                           AAX14711;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               bacteria.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                                                                                                                                                                     RESULT 2507
AAX14711
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This sequence is an immunosuppressant inhibitor oligonucleotide, which is used in the invention. The invention relates to a composition which contains at least one inhibitor (less than 100 kD) of a substance (e.g. transforming growth factor TGF-beta, vascular endothelial growth factor TGF-beta, vascular endothelial growth factor VEGF, interleukin-10 IL-10, prostaglandin E2 PGE2, or their receptors) that adversely affects the immune response. The composition also includes at least one stimulant that positively affects the immune response. This oligonuclectide is an example of an inhibitor that is used in the composition. The composition is used as an immunostimulant for the composition. The composition is used as an immunostimulant for the composition, stomach, infections, particularly hyperproliferation, leukaemia, (non-)Hodgkin's lymphona; carcinoma (of oesophagus, bronchi, colon-rectum, stomach, intestine, gall bladder or duct, pancreas, anus, maignant melanoma, brain tumours and sarcomas. The oligonucleotides, most of which are directed against TGFbeta or VEGF, are inhibitors of monocyte chemotactic protein.
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0
                                                                                                                                                                           Immunosuppressant inhibitor; transforming growth factor beta; TGF beta; vascular endothelial growth factor; VEGF; interleukin-10; IL-10; cancer; prostaglandin E2; PGE2; immune response; tumour; asthma; Crohn's disease; glomerulonephritis; protein-1; MCP-1; ulcerative colitis; diabetes; atherosclerosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      collis, quabetes, glomerulonephritis, acute respiratory distress syndrome and the formation of atherosclerotic plaque
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Composition containing immune stimulant and inhibitor of agent adversely affects the immune response, for treating cancers and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
                                                                                                                                               Immunosuppressant inhibitor oligonucleotide TGF-beta-3-rwk-14.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12.9%; Score 9.4; DB 1; Length 14; 90.9%; Pred. No. 1.4e+03; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   inflammatories for treating e.g. asthma, Crohn's disease, colitis, diabetes, glomerulonephritis, acute respiratory o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 14 BP; 1 A; 7 C; 0 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (BIOG-) BIOGNOSTIK GES BIOMOLEKULARE DIAGNOSTIK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Schlingensiepen K, Schlingensiepen R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 10; Fig 1; 30pp; English.
                                   BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                         99WO-EP004013.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               98EP-00110709.
                                   AAZ65640 standard; DNA; 14
                                                                                                                (first entry)
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Best Local Similarity 90.9
Matches 10; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          932 CCCTCCTCTTC 942
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                                                                                                                30-MAR-2000
                                                                                                                                                                                                                                                                                                                                                                           409963975-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       10-JUN-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       25-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                   16-DEC-1999,
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                                                                            AAZ65640;
RESULT 2508
                     AAZ65640
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Gaps

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Local Similarity 90.5 nes 10; Conservative

Matches

931 TCCCTCCTCTT 941

schultz1-899.rng

cell expresses telomerase activity and its RNA component

Sequence 14 BP; 8 A; 2 C; 2 G; 2 T; 0 U; 0 Other;

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The present sequence represents a peptide nucleic acid molecule which hybridises to the mRNA component of mammalian telomerase, and inhibits celomerase activity. Telomerase is a ribonucleoprotein enzyme that synthesizes one strand of the telomeric DNA, using as a template an 11 nucleotide sequence contained within the RNA component of the enzyme. The nucleotide sequence contained within the RNA component of the enzyme. The nucleotide sequence contained within the RNA component of the enzyme. The nucleotide sequence contained hybrid acquired for once than 25 bases, which include the sequence of TYAGG. The uncharged nature of the PNA backbone increases the melting temperature of associating strands, affords greater resistance of degradation by proteases or nucleases. The concerns may be used for treating disease or nucleases. The therapeutic PNNs may be used for treating disease ornditions such as cancers, neoplasia, hyperplasia, neurodegenerative diseases. The immunodeficiency virus (HIV) infection/AIDS (acquired immunodeficiency virus (HIV) infection and other rucleotides PNAs may be used for individuals, e.g. paternity testing, based on hrs gene restriction of individuals, e.g. paternity testing, based on hrs gene restriction allows cancerous conditions to be detected with increased confidence and proper or detect of passibly at an earlier stage, before cells are detected as cancerous conditions to be detected with increased confidence of the present invention central or methods of the present invention central and proper and immort 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New peptide nucleic acid (PNA) compounds that inhibit telomerase activity in mammalian cells is useful as probes to detect the RNA component of a mammalian telomerase.
                                                                                                                                                                                                                                                                                                                                                                                                                                  Peptide nucleic acid, PNA, telomerase, ribonucleoprotein enzyme, cancer, inhibitor, neoplasia, neurodegenerative disease, aging, hyperplasia, AIDS, HIV, fungal infection, forensic identification, detect, tumour,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /*tag= a // hote= "Peptide nucleic acid molecule, where N-(2-aminoethyl)glycine units are linked to nucleotide bases via glycine amino N through a methylenecarbonyl linker"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Wright WE, Piatyszek MA, Shay JW, Norton JC, Corey DR;
                                                                                                                                                                                                                                                                                                                                            PNA sequence #50 used to inhibit telomerase activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cocation/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 2; Col 37; 45pp; English.
                                                   AAA37592 standard; DNA; 14 BP.
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                                                                                                                                                                                                                                                (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       paternity testing; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2000-292432/25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Key
misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       09-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   09-APR-1996;
                                                                                                                                                                                                                                                15-AUG-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Synthetic.
                                                                                                                                               AAA37592;
AAA37592,
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The invention relates to the regulation of adenoviral packaging. The method of the invention comprises propagating an adenoviral vector containing a repressor binding site, in the absence of the repressor. After propagation, vector packaging is repressed by the appropriate repressor protein. The invention also encompasses an adenoviral vector chicken ovalbumin upstream promoter transcription factor) binding sites (hAZ59919). Adenoviral vectors containing sepressor binding sites are used for DNA delivery, e.g., for expression of a therapeutic protein; in genetic immunisation; or to produce antiviral DNA or antisense RNA. Typical heterologous genes that can be expressed include those for interleukin-2, alphal-antitrypsin, cystic fibrosis transmembrane conductance regulator and coagulation factor VIII. These vectors have very large capacity (up to 36 kp) for foreign DNA and minimise the risk of generating replication competent virus (since vector and helper virus can be designed such that they have no overlapping packaging sequences that might permit homologous recombination). The presence of the repressor binding site allows selective inhibition of virion production (i.e., packaging of ne vector in presence of another). Sequences AZ59890-Z59896 represent adenovirus minimal packaging elements.
                                    ö
                                                                                                                                                                                                                                                                                                                              Adenovirus; minimal packaging element; A repeat; repressor binding site;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Regulating adenoviral packaging by incorporation of repressor binding sites that allow selective suppression of packaging, used for gene
                                      Gaps
                                    .
0
   Length 14;
                                  1; Indels
                                                                                                                                                                                                                                                                                             Adenovirus minimal packaging element, A repeat AII.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 14 BP; 1 A; 3 C; 3 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Schmid SI, Ostapchuk PH, Erturk E;
 9.4; DB 1;
No. 1.4e+03;
                                  0; Mismatches
                   Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (UYNY ) UNIV NEW YORK STATE RES FOUND.
   Score
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Page 15; 71pp; English.
                                                                                                                                                                                           AAZS9891 standard; DNA; 14 BP.
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 12.9%;
90.9%;
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Query Match
Best Local Similarity 90.9
Matches 10; Conservative
                                                                     910 TICTITGGICT 920
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                                                                                                                                                                                                                                                                                                                                                  DNA delivery; ds
                                                                                                                                                                                                                                                                                                                                                                                  Mastadenovirus.
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                                                                                                                                                                                                                                                              08-MAY-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hearing P,
                                                                                                                                                                                                                           AAZ59891;
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                                                                                                                                                                         AAZ59891
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/*tag= a
/note= "This sequence is a peptide nucleic acid, i.e. it
contains a polyamide backbone instead of a deoxyribose
backbone"
                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New peptide nucleic acids that hybridizes to the RNA component of mammalian telomerase, useful for treating or preventing cancer, inflammation, lymphoproliferative diseases, autoimmune disease, or neurodegenerative diseases.
                                                                                                                                                                                                                       Mammalian; peptide nucleic acid; probe; forensic; paternity testir human telomerase RNA component; hTR gene RFLP pattern; cancer; inflammation; lymphoproliferative disease; autoimmune disease; neurodegenerative disease; nopplasia; hyperplasia; HTV; AIDS; human immunodeficiency virus; acquired immunodeficiency syndrome;
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                                                                                                                                                                                                                                                                                 telomere metabolism; mutant; cytostatic; anti-inflammatory; immunosuppressive; polyamide backbone; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Norton JC;
                                                                                                                                                                                                 PNA 1 inhibiting human and mammalian telomerase activity.
                        1; Indels
Score 9.4; DB 1;
Pred. No. 1.4e+03;
0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Corey DR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 2; Col 37-38; 46pp; English.
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                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                     .463/c
AAS15463 standard; DNA; 14 BP.
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97US-00838545.
 12.9%;
90.9%;
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                          10; Conservative
                                               902 TGGTCATTTTC 912
                                                                      TGGCCATTITC 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Shay JW, Wright WE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2001-638024/73.
Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                      Key
modified_base
                                                                                                                                                                                                                                                                                                                   Homo sapiens.
Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  08-JUL-1999;
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09-APR-1997;
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                                                                                                                                                      AAS15463
                        Matches
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The invention relates to a polymerase chain reaction (PCR) based method of DNA fingerprinting, comprising using primers that match the conserved regions of a gene family. The method is useful for gene expression analysis of any cell or tissue, or for the performance of DNA fingerprinting analysis of the same organism in order that one will reveal the function of a gene that produced differential product between genotypes. The method is also useful for identifying pCR reactions that contain a gene of interest in a gene family reverse transcriptase (RT) PCR expression analysis. The method is also useful for identifying genes that to a gene family that might be involved in cancer formation. The method is particularly useful for comparing genomic sequences. These are also applicable in agriculture (e.g. to mark useful genes to assist breeding). The current sequence represents an animal cis-regulatory sequence. This is used in DNA fingerprinting using primers or a mix of primers that match the sequence of ubiquitous cis-acting regulatory elements. (Updated on 29-AUG-2003 to standardise OS field)
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                                                           the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Polymerase chain reaction based method of DNA fingerprinting, useful for analyzing genes, e.g. for identifying genes involved in cancer formation, involves using a mix of primers that match the conserved regions of a
hyperplasia, human immunodeficiency virus (HIV) infections, acquired immunodeficiency syndrome (AIDS) and associated pathologies, and other diseases characterised by abnormal telomere metabolism or telomerase
                                                         activity. The present sequence represents one of the PNA sequences of
                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DNA fingerprinting, cancer; agriculture; breeding; PCR; primer; gene family; ds.
                                                                                                                                                                                   ;
0
                                                                                                                                         Score 9.4; DB 1; Length 14;
Pred. No. 1.46+03;
0; Mismatches 1; Indels
                                                                                                              2 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (GENE-) GENENA LTD.
(AGRI-) AGRIC RES ORG NEWE YA'AR RES CENTE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Animal cis-regulatory sequence from MyoD.
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02-JUL-2000; 2000IL-00137124.
20-AUG-2000; 2000IL-00137959.
                                                                                                                                                 12.9%;
                                                                                                              Sequence 14 BP; 8 A; 2 C; 2
                                                                                                                                                                                                                                                                                                                                                    ABL42252 standard; DNA; 14
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(first entry)
                                                                                                                                                                                     10; Conservative
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                                                                                                                                               Query Match
Best Local Similarity
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01-JUL-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         gene family.
                                                                                invention
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comprising

The present invention relates to peptide nucleic acids (PNAs), comprising as sequence of 6-25 nucleobases, that inhibit telomerase activity in mammalian cells by hybridising to the RNA component of mammalian telomerase and sequence as probes to detect the RNA component of mammalian telomerase and as inhibitors of telomerase activity, or to detect and/or quantitate polynuclectide having the human telomerase RNA component (hTR) sequence, as well as in forensic identification of individuals, such as parently testing or identification of suspects or unknown descendants based on the hTR gene RPLP pattern. The PNA can be further used for treating or preventing cancer, inflammation, hymphoproliferative diseases, autofimmune disease, or neurodegenerative diseases. The PNAs in combination with other pharmaceuticals (such as antineoplastic or cytostatic agents) can be used for treating neoplasia,

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The antisense oligonucleotides are useful in the treatment of tumours in which expression of TGF-beta is of relevance for pathogenicity and/or inhibition of pathological angiogenesis. They are used especially for the treatment of the immunosuppressive effect of TGF-beta, augmentation of the proliferation of cytotoxic lymphocytes, treatment of endogenous hyperexpression of TGF-beta, treatment of breast tumours, neurofibromas and mallignant glindmas, including glioblastomas, treatment and prophylaxis of skin carcinogenesis, and treatment of oesophageal and gastric carcinomas. See AAQ78352-078488 are antisense oligodeoxynucleotides of TGF-beta 2 in the form of phosphorothioate analogues. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          H-ras; wild-type; immobilising; diagnosis; ethylene acrylic acid; ethylene methacrylic acid; polypropylene; biotin; cystic fibrosis; array; ss.
                                                                                                       Transforming growth factor beta; TGF-beta; antisense; treatment; tumour; angiogenesis; breast tumour; neurofibroma; glioma; glioblastoma; carcinogenesis; carcinoma; oesophagus; oesophageal; gastric; gut; immunosuppression; oligonucleotide; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New transforming growth factor beta anti:sense oligo:nucleotide(s) treating immunosuppression, tumours, etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Schlingensiepen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          12.6%; Score 9.2; DB 1; Length 14; 78.6%; Pred. No. 1.5e+03; attive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          One from an array of 58 cystic fibrosis oligonucleotides.
                                                                    TGF-beta gene phosphorothioate antisense oligonucleotide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Schlingensiepen K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 14 BP; 1 A; 2 C; 5 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                    (BIOG-) BIOGNOSTIK GES BIOMOLEKULARE DIAGNOSTIK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 6; Page 58; 74pp; English.
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93EP-00107849.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAV06882 standard; DNA; 14
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                (revised)
(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         Schlingensiepen G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
Matches 11; Conserv
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                                                                                                                                                                                                                                                                                                                         29-APR-1994;
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                  25-MAR-2003
27-JUN-1995
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                                                                                                                                                                                                          Synthetic.
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                                                                                             Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA coding for human factor IC - used for producing polypeptide and detecting genetic modifications in diagnosing blood clotting
                                                                                                                                                                                                                                                                                                                                                                                                                      Human factor IX; genetic deficiencies; blood clotting disorders;
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Best Local Similarity 64.3%; Pred. No. 1.5e+03;
Matches 9; Conservative 3; Mismatches 2; Indels
                                                    Query Match 12.9%; Score 9.4; DB 1; Length 14; Best Local Similarity 90.9%; Pred. No. 1.46+03; Matches 10; Conservative 0; Mismatches 1; Indels
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                  Sequence 14 BP; 4 A; 4 C; 4 G; 2 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                     BP.
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86US-00888041.
87US-00094031.
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                                                                                                                                                                                                                                                                   AAQ10579 standard; DNA; 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAQ78469 standard; DNA; 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                           haemophilia B; ss
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
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18-JUL-1986;
28-AUG-1987;
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AAQ78469

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Claim 10; Page 10; 15pp; Japanese.
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Local Similarity 78.6%;
tes 11; Conservative (
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                                                                                                                                                                                                                                             13-AUG-1998 (first entry)
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                                                                                                        11; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  anti-metastatic agent
                                                                                            Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                             (TERU ) TERUMO CORP.
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                                                                                                                                                                                                                                                                                                                                          JP10127286-A.
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                                                                                                                                                                                                                                                                                        Hepatocyte
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                                                                                   Query Match
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AC AAV9
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DT 01-M
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                                                                                                                                                                                                                      The present sequence represents one of an array of 58 cystic fibrosis objogonaclectides. The invention relates to a new reagent for immobilising a biopolymer. It comprises a solid support fabricated from a polymeric material having at least one surface comprising pendant acyl fluoride functionalities. The reagent is stable under conditions for synthesising and immobilising biopolymers and is stable under conditions for synthesising analyse the biopolymers. The reagents can be formed into devices which are physically rugged and inexpensive which can be used in analytical and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Oligo:nucleotide inhibiting HGF production - useful as antitumour and
                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                 Polymeric reagents for immobilising biopolymers - are stable under synthesis conditions.
                                                                                                                                                                                                                                                                                                                                                                            ..
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hepatocyte growth factor; HGF; c-Met; modulator; inhibitor; antitumour agent; anti-metastasis agent; primer; ss.
                                                                                                                                                                                                                                                                                                                                                   Query Match 12.6%; Score 9.2; DB 1; Length 14; Best Local Similarity 78.6%; Pred. No. 1.5e+03; Matches 11; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hepatocyte growth factor inhibiting oligonucleotide #17.
                                                                                                                                                                                                                                                                                                                                 Sequence 14 BP; 1 A; 3 C; 3 G; 7 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 10; Page 10; 15pp; Japanese.
                                                                                                                                                                                                    Example 7; Fig 19; 66pp; English
                                                              97WO-US008880
                                                                                  96US-00658664
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                                                                                                     (BECI ) BECKMAN INSTR
                                                                                                                                                                                                                                                                                                            diagnostic procedures
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                    WO9746597-A1
                                                              22-MAY-1997;
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                                         11-DEC-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19-MAY-1998.
 Synthetic
                                                                                                                           Milton RC;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAV11925
                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 2516
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AAV11909-V11925, AAV11927 and AAV11928 are oligonucleotide primers used to identify sequences which modulate or inhibit expression, production or reception of hepatocyte growth factor (HGF) or expression of c-Met. Such oligonucleotides are useful as antitumour or anti-metastasis agents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAVI1909-V11925, AAVI1927 and AAVI1928 are oligonucleotide primers used to identify sequences which modulate or inhibit expression, production or reception of hepatocyte growth factor (HGF) or expression of G-Met. Such oligonucleotides are useful as antitumour or anti-metastasis agents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Oligo:nucleotide inhibiting HGF production - useful as antitumour and
                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              growth factor; HGF; c-Met; modulator; inhibitor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 9.2; DB 1; Length 14;
Pred. No. 1.5e+03;
0; Mismatches 3; Indels
                                                                                                                                                                                                                                                               12.6%; Score 9.2; DB 1; Length 14; 78.6%; Pred. No. 1.5e+03; ive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hepatocyte growth factor inhibiting oligonucleotide #16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              antitumour agent; anti-metastasis agent; primer; ss.
                                                                                                                                                                                          Sequence 14 BP; 0 A; 8 C; 0 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 14 BP; 6 A; 0 C; 8 G; 0 T; 0 U; 0 Other;
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Fresco JR;
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                                                                                                                                                                                                                                     WPI; 1999-327425/27.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              10-NOV-1998;
                                         Homo sapiens.
                                                                  WO9924622-A1.
                                                                                                                          10-NOV-1998;
                                                                                                                                                     10-NOV-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO9924622-A1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20-MAY-1999.
                                                                                                                                                                                                          Johnson MD,
                                                                                              20-MAY-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAX61148;
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                                                                                                                                                                                                                                                                                                                                                                                                    The present invention describes enzymatic nucleic acid molecules with RNA cleaving activity (e.g. ribozymes) which are capable of modulating the expression of plant genes: (i) involved in biosynthesis of alkaloids; or (ii) involved in flower formation. AAV95802 to AAV96334, and AAV96335 to AAV96354 represent potato solanidine glucosyltransferase hammerhead and AAV96734 represent potato solanidine glucosyltransferase target sequences. AAV96773 to AAV97170, and AAV97171 to AAV97195 represent potato colanidine glucosyltransferase target sequences. AAV96773, and AAV97170, and AAV97171 to AAV97195 represent colarate synthase target sequences. Ribozymes of the present invention can be used to inhibit the synthesis of toxic alkaloids in solanaecous plants, particularly potato but also tomato, pepper, aubergine and ditura or to inhibit flowering in potato, lettuce, spinach, cabbage, brussel sprouts, arugula, kale, collards, chard, beet, turnip, sweet potato and turf grass. Also the ribozymes can be used for RNA manipulation in the same way that restriction endonucleases are for DNA, as well as to examine ribozymes can be targeted to specific genes or to consensus sequences within a family of related genes, and being catalytic need to be present at only very low concentrations
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                     New enzymatic nucleic acid(s) - useful for, e.g. reducing alkaloid biosynthesis or regulating flowering.
                          Solanidine, glucosyltransferase, potato, citrate synthase, target, hammerhead ribozyme, hairpin ribozyme, alkaloid biosynthesis; flower formation, cleavage, solanaceous plant, ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Probe; human; chromosome 17 triple-helix forming oligonucleotide;
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  Potato citrate synthase target sequence position 539.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 14 BP; 3 A; 5 C; 1 G; 0 T; 5 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human chromosome alpha-satellite region.
                                                                                                                                                                                                                                                                                                                                                                                Claim 54; Page 59; 79pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAX61182 standard; DNA; 14 BP.
                                                                                                                                                                                             97US-0036545P.
97US-0036599P.
97US-00979416.
                                                                                                                                                                     98WO-US000738
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                                                                                                                                                                                                                                                     (RIBO-) RIBOZYME PHARM INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 UCCCUGAUCAUCAU 14
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                                                                                                                                                                                                                                                                                  Zwick MG, Mcswiggen JA;
                                                                                                                                                                                                                                                                                                            WPI; 1998-427939/36.
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Best Local Similarity
Matches 6; Conserv
                                                                                    Solanum tuberosum
                                                                                                               WO9832843-A2
                                                                                                                                                                     14-JAN-1998;
                                                                                                                                                                                                28-JAN-1997;
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                                                                                                                                         30-JUL-1998
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This sequence represents a human chromosome alpha-satellite region. The invention relates to the use of a triple-helix forming oligonucleotide for in situ detection of a double-stranded target nucleic acid sequence. The method can be used to detect a genetic disorder e.g. to detect an extra or missing chromosome or fragment or aneuploidy, especially for be used to screen for individuals at risk of developing a disease or for disgnosing an infectious disease. The use of triple helix forming of jigonucleotides allows in situ detection of double stranded target sequence as opposed to prior art uses of developing potential anti-gene therapeutic agents or artificial restriction endonucleases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel use of triple helix forming oligonucleotides, useful for in situ detection of double stranded target sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Probe; human; chromosome 17 triple-helix forming oligonucleotide; genetic disorder; missing chromosome; aneuploidy; chromosome 21; infectious disease; diagnosis; alpha-satellite region; ss.
genetic disorder; missing chromosome; aneuploidy; chromosome 21; infectious disease; diagnosis; alpha-satellite region; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 14 BP; 7 A; 0 C; 6 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human chromosome alpha-satellite region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 19; Page 13; 45pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAX61148 standard; DNA; 14 BP
                                                                                                                                                                                                                                                                            97US-0064997P.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TTTCCCTTTTCACC 1
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This sequence represents a human chromosome alpha-satellite region. The invention relates to the use of a triple-helix forming oligonuclectide for in situ detection of a double-stranded target nucleic acid sequence. The method can be used to detect a genetic disorder e.g. to detect an extra or missing chromosome or fragment or aneuploidy, especially for detecting an extra or missing chromosome 17 or 21. The method can be also be used to screen for individuals at risk of developing a disease or for diagnosing an infectious disease. The use of triple helix forming oligonuclectides allows in situ detection of double stranded target sequence as opposed to prior art uses of developing potential anti-gene therapeutic agents or artificial restriction endonucleases
                                                                                  triple helix forming oligonucleotides, useful for in situ double stranded target sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                12.6%; Score 9.2; DB 1; Length 14; 78.6%; Pred. No. 1.5e+03; tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 14 BP; 8 A; 0 C; 5 G; 1 T; 0 U; 0 Other;
                                                                                                                                                Claim 19; Page 11; 45pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       921 ITGCCTTTTATCCC 934
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAX14931 standard; DNA; 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 70.0.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                14 rrccrrrrcrace 1
  Johnson MD, Fresco JR;
                                         WPI; 1999-327425/27.
                                                                                  Novel use of
detection of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAX14931;
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Triplex formation; DNA detection; triple helix; identification; bacteria;
                                Triple helix third strand of 23S rRNA gene nucleotides 663-676.
                        (first entry)
                                            oncogene; virus; ss.
                        24-MAR-1999
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Synthetic. Haemophilus influenzae.

US5861244-A.

19-JAN-1999,

93US-00173489,

(PROF-) PROFILE DIAGNOSTIC SCI INC.

92US-00968436.

29-OCT-1992;

Wang C; Hepburn AG,

WPI; 1999-130384/11.

Assay of genetic sequences based on triplex formation from double stranded analyte - and hybrid of anchor and reporter sequences, with reporter released if triplex formation occurs, used e.g. to identify bacteria.

The present sequence represents a potential triple-helix forming region. It can be used to demonstrate the assay of the invention. The assay occuprises adding a sample containing double-stranded DNA test sequences, e.g. containing the present sequence, to an aqueous medium containing at least one complex of anchor DNA, attached to a solid support, and reporter DNA, where either a part of the anchor DNA or reporter DNA is designed to form a triple-strand structure with part of the test sequence. Triplex formation results in displacement of the reporter DNA which is detected as an indication of the presence of the DNA test sequence. The method is used to detect DNA sequences, particularly for identification of bacteria go detecting genes for ribosomal RNA) in clinical samples, but also detection of oncogenes and Hepatitis B virus

Sequence 14 BP; 8 A; 0 C; 6 G; 0 T; 0 U; 0 Other;

Disclosure; Col 23-24; 168pp; English.

The present sequence represents a polynucleotide that is able to form a triple helix with a double stranded sequence. Cytosine bases in the present can be replaced with 5-methylcytosine for increased triplex stability. The present sequence is used in the assay of the invention,

ö assay comprises adding a sample containing double-stranded DNA test sequences to an aqueous medium containing at least one complex of anchor DNA, attached to a solid support, and reporter DNA, where either a part of the anchor DNA or reporter DNA is designed to form a triple-strand structure with part of the test sequence. Triplex formation results in displacement of the reporter DNA which is detected as an indication of the presence of the DNA test sequence. The method is used to detect DNA sequences, particularly for identification of bacteria (by detecting genes for ribosomal RNA) in clinical samples, but also detection of oncogenes and Hepatitis B virus can be part of the anchor DNA or reporter DNA sequence. The Gaps Assay of genetic sequences based on triplex formation from double stranded analyte - and hybrid of anchor and reporter sequences, with reporter released if triplex formation occurs, used e.g. to identify ; Triple-helix forming region; Triplex formation; DNA detection; identification; bacteria; oncogene; virus; ds. Triple helix forming nucleotides 250-263 of Prealbumin gene. Score 9.2; DB 1; Length 14; Pred. No. 1.5e+03; 3; Indels Sequence 14 BP; 0 A; 8 C; 1 G; 5 T; 0 U; 0 Other; 0; Mismatches Disclosure, Col 17-18; 168pp, English. (PROF-) PROFILE DIAGNOSTIC SCI INC. AAX14710 standard; DNA; 14 BP. 12.6%; 78.6%; 93US-00173489. 92US-00968436. 931 TCCCTCCTCTTCAT 944 1 rccrccccarcri 14 (first entry) 11; Conservative Hepburn AG, Wang C; WPI; 1999-130384/11. Query Match Best Local Similarity 22-DEC-1993; 24-MAR-1999 29-OCT-1992; Homo sapiens 19-JAN-1999. US5861244-A. AAX14710; where it bacteria. Matches RESULT 88888888888888 ઠે g

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Gaps

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AAD07946 standard; DNA; 14 BP.

RESULT 2524 AAD07946

AAD07946;

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The present sequence represents a polynucleotide that is able to form a triple helix with a double stranded sequence. Cytosine bases in the present can be replaced with 5-methylcytosine for increased triplex tability. The present sequence is used in the assay of the invention, where it can be part of the anchor DNA or reporter DNA sequence. The assay comprises adding a sample containing double-stranded DNA test DNA, attached to a solid support, and reporter DNA, where either a part of the anchor DNA or reporter DNA, where either a part of the anchor DNA or reporter DNA, where either a part of the nathor DNA or reporter DNA, where either a part of the part of the test sequence. Triplex formation results in displacement of the reporter DNA which is detected as an indication of the presence of the DNA test sequence. The method is used to detect DNA sequences, particularly for identification of bacteria (by detecting genes for ribosomal RNA) in clinical samples, but also detecting oncogenes and Hepatitis B virus
                                    ö
                                                                                                                                                                                                                                                                                                                                               Triplex formation; DNA detection; triple helix; identification; bacteria;
                                    Gaps
                                                                                                                                                                                                                                                                                                            Triple helix third strand of retinoblastoma gene nucleotides 281-394.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Assay of genetic sequences based on triplex formation from double stranded analyte - and hybrid of anchor and reporter sequences, with reporter released if triplex formation occurs, used e.g. to identify bacteria.
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Query Match
12.6%; Score 9.2; DB 1; Length 14;
Best Local Similarity 78.6%; Pred. No. 1.5e+03;
Matches 11; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 14 BP; 0 A; 3 C; 0 G; 11 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Col 15-16; 168pp; English.
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                                                                         926 TITIAICCCICCIC 939
                                                                                                                                                                                                    AAX14691 standard; DNA; 14
                                                                                                                                                                                                                                                                            (first entry)
                                                                                                        14 rrrrrccrccrc
                                                                                                                                                                                                                                                                                                                                                                      oncogene; virus; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Wang C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1999-130384/11.
                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    22-DEC-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       29-OCT-1992;
                                                                                                                                                                                                                                                                            24-MAR-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hepburn AG,
                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic.
                                                                                                                                                                                                                                         AAX14691;
                                                                                                                                                                RESULT 2523
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                                                                                                                            Human; antisense; amyloid precursor protein; APP; amyloid beta protein; AbetaP; Alzheimer's disease; cognitive ability; antisense therapy; nootropic; neuroprotective; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel antisense compounds for modulating expression of amyloid beta protein in cells or tissues and for preventing, treating conditions associated with expression of amyloid beta protein, e.g. Alzheimer's
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 14 BP; 5 A; 6 C; 0 G; 3 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RNA oligonucleotide #2 used in a binding assay.
                                                                                               OL-3
                                                                                               Human antisense oligonucleotide,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 10; Page 6; 70pp; English.
                                                                                                                                                                                                                                                                                                                                        99US-00458481
                                                                                                                                                                                                                                                                                                        08-DEC-2000; 2000WO-US03383
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        oligonucleotide
                                                                                                                                                                                                                                     WO200142266-A1.
                                                                                                                                                                                                                                                                                                                                          09-DEC-1999;
                                                                                                                                                                                                    Homo sapiens
                                                                 06-AUG-2001
                                                                                                                                                                                                                                                                       14-JUN-2001,
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                                                                                                                                                                                                                                                                                                                                                                                                           Kumar VB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
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Gaps

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Query Match
12.6%; Score 9.2; DB 1; Length 14;
Best Local Similarity 78.6%; Pred. No. 1.5e+03;
Matches 11; Conservative 0; Mismatches 3; Indels

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The invention comprises a method of manufacturing a novel DNA chip (uni chip), using reverse transcriptase. The invention further comprises a method of detecting novel genes (using the novel DNA chip). The manufacturing method comprises the steps of: preparing various kinds of primers on a DNA chip by annealing an oligonucleotide primer having a specific sequence to a DNA chip having a poly T tail; complementarily annealing unsequenced mRNA to the primers; adding reverse transcriptase to synthesize cDNA on the DNA chip; and removing mRNA therefrom using RNAse to obtain a cDNA library chip having only cDNA
                                                             enzyme to detect
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /*tag= a
/bound_moiety= "Ribozyme complex strand #1"
/note= "Binds to nucleotides 14-11 of the RNA sequence
shown in (AAL50499)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ribozyme complex RNA strand; RNA structural properties; IP-RP-HPLC; ion pairing reverse phase high performance liquid chromatography; ss; intramolecular interaction; three-dimensional structure.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Analyzing RNA by partially hydrolyzing RNA, separating and detecting cleaved RNA by high performance liquid chromatography, and absence of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /*tag= b
/bound moiety= "Ribozyme complex strand #1"
/note="Binds to nucleotides 6-1 of the RNA sequence
shown in (AALS0499)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      .
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ouery Match 12.6%; Score 9.2; DB 1; Length 14; Best Local Similarity 78.6%; Pred. No. 1.5e+03; Matches 11; Conservative 0; Mismatches 3; Indels
                                                       Manufacturing of DNA chip using reverse transcriptase novel genes comprises genetic recombinant techniques.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 14 BP; 1 A; 2 C; 1 G; 10 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                            Disclosure; Page 5; 6pp; Korean.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ribozyme complex RNA strand #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 29-NOV-2000; 2000US-00727138.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TTTTCTTTGGTCTT 921
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                rrrrrragcrcrr 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAL50500 standard; RNA; 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             9. .14
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     WPI; 2002-301918/34.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4PI; 2002-690387/74.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (HORN/) HORNBY D P
(DICK/) DICKMAN M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 JS2002094539-A1
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misc_binding
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAL50500;
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        EXETXXXXCCCCCCCXXX
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present invention relates to an oligomer comprising L-ribo-
configurated Locked Nucleoside Analogues (L-ribo-LNA analogues). The
present sequence is an RNA oligomuclectide. Binding studies of the L-ribo
-LNA analogues towards the present sequence were carried out, to
determine the thermostability of the L-ribo-LNA analogues. The analogue of
the present invention have a variety of uses e.g. in the preparation of
conjugates of the L-ribo-LNA modified oligonucleotides (oligomers)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Oligomers comprising L-ribo-Locked Nucleic Acid (LNA) nucleosides, useful for therapeutic purposes e.g. in the construction of oligonucleotides, as substrates for nucleic acids polymerases and in RNA mediated catalytic
L-ribo-configurated Locked Nucleoside Analogue; L-ribo-LNA analogue; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Novel DNA chip; ss; manufacture; uni chip; reverse transcriptase; novel gene detection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 9.2; DB 1; Length 14; Pred. No. 1.5e+03; 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 14 BP; 13 A; 1 C; 0 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 11; Page 56; 79pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BP.
                                                                                                                                                                                                                                                                            04-MAY-1999; 99DK-0000603.
01-SEP-1999; 99DK-00001225.
11-JAN-2000; 2000DK-00000032.
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78.6%;
                                                                                                                                                                                                                        04-MAY-2000; 2000WO-DK000225
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAL42800 standard; DNA; 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITITIGITITI 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 78.6
Matches 11; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                      (EXIQ-) EXIQON AS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (SONG/) SONG K H.
                                                                                                         WO200066604-A2.
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                                                       Unidentified
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Unidentified
                                                                                                                                                                   09-NOV-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                             Wengel J;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               14
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AAL42800
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AC AAL42800
AC AAL4280
DT 05-AUG-;
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DX Novel DI
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                                                                                            The invention comprises a method for analysing the structural properties of an RNA molecule. The method of the invention involves contacting the RNA molecule with a cleaved reagent capable of partially hydrolysing the RNA. The cleaved RNA is then separated and detected by ion pairing reverse phase high performance liquid chromatography (IP-RP-HPLC) - absence of cleavage events in a region of the RNA indicates that the region is relatively inaccessible to solvent. The method of the invention is useful for analysing the structural properties of the RNA molecule, including region(s) that are relatively inaccessible to solvent owing to intramolecular interactions. The method is used to characterise the three dimensional structure of an RNA molecule, and is used to characterise the interaction of an RNA (e.g. a ribozyme) with its substrate, where the intermolecular interaction is between the RNA molecule and an RNA binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present invention discloses seven kinds of full length and partial sequences of a yak milk protein gene. They include alpha-lactoalbumin gene full length sequence, alpha-lactoalbumin gene 5' lateral wing and 3' terminal sequence, beta-lactoglobulin gene 5' lateral wing and 3' terminal sequence, alpha S1-casein gene 5' lateral wing and 3' terminal sequence, alpha S2-casein gene 5' lateral wing and 1' terminal sequence, lateral wing and 3' terminal sequence, alpha S2-casein gene 5' alteral wing and 3' terminal sequence, kappa casein gene 5' lateral wing and 3' terminal sequence, and lactoferritin gene 5' lateral wing
                                                                                                                                                                                                                                                                                                                                                                            protein. The present RNA sequence represents a ribozyme complex strand that was used in an example of the invention
cleavage in region of RNA indicates that the region is inaccessible to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               yak milk; alpha-lactoalbumin; beta-lactoglobulin; alpha S1-casein; alpha S2-casein; beta-casein; kappa-casein; lactoferritin; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12.6%; Score 9.2; DB 1; Length 14; 78.6%; Pred. No. 1.5e+03; tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                          Seguence 14 BP; 8 A; 1 C; 4 G; 0 T; 1 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 8 (disclosure); 41pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Seven kinds of yak milk protein gene sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Yak milk protein gene related oligo, 454-467.
                                                             Example 2; Fig 2; 16pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADE64664 standard; DNA; 14 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  08-DEC-2000; 2000CN-00134189
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 2002-741796/81.
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bos grunniens
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ADE64664
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human, gene therapy, adenosine deaminase deficiency, p53, beta-globin, retinoblastoma, BRCA1, BRCA2, CFTR, cystic fibrosis, cancer, Factor Vy, cyclin-dependent kinase inhibitor 2A, CDKN2A, melanoma, AEC, HBA1, HBA2, adenomatous polyposis of the colon, Factor VII, Factor IX, thrombosis; haemophilia, alpha thalassaemia, haemoglobin alpha locus 1, MLH1, APOE, mismatch repair, MSH2, MSH6, hyperlipidaemia, apolipoprotein E, LDLR, familial hypercholesterolaemia, UGT1, syndrome, APP, PSEN1, antisense; UDP-glucuronosyltransferase, amyloid precursor protein; presentilin-1, Albehmer's disease, cytostatic, antisickling; antianaemic, haemostatic;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Oligonucleotide for targeted alterations of genetic sequences and for
treating cystic fibrosis, comprises at least one mismatch and chemical
                                                                                                                                  Gaps
sequence. This polynucleotide sequence represents an oligo relating the yak milk protein genes of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                      Retinoblastoma mutation correcting oligonucleotide SEQ ID NO: 560.
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                                                                                            Length 14;
                                                                                         Score 9.2; DB 1; Length 14
Pred. No. 1.5e+03;
0; Mismatches 3; Indels
                                                       Sequence 14 BP; 1 A; 3 C; 1 G; 9 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Rice MC;
                                                                                                                                                                                                                                                                                            714/c
ABA77714 standard; DNA; 17 BP.
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27-MAR-2000; 2000US-0192179P.
01-UUN-2000; 2000US-0208538P.
30-OCT-2000; 2000US-0244989P.
                                                                                            Query Match
Best Local Similarity 78.6%;
Matches 11; Conservative (
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                                                                                                                                                                         937 CTCTTCATTGGTTT 950
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1 CACTTCTTTTGTTT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (UYDE ) UNIV DELAWARE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           antilipemic; ss
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modification.
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Gaps

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Indels

0; Mismatches

Score 9.2; DB 1; Length 17; Pred, No. 1.6e+03;

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haemophilia, hypercholesterolaemia, thalassaemia, sickle cell anaemia, Alzheimer's disease, melanoma, adenomatous polyposis of the colon and various syndromes. The present sequence is one of the gene correcting oligonucleotides of the invention
                                                                                                                          Sequence 17 BP; 6 A; 2 C; 4 G; 5 T; 0 U; 0 Other;
                                                                                                                                                                        12.6%;
78.6%;
                                                                                                                                                                                                                         11; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (revised)
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                                                                                                                                                                                                  Best Local Similarity
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21-FEB-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO9521912-A1
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20-MAR-1996
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Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             be used for the targeted alteration of genomic sequences, where the oligonuclectide has at least one mismatch compared with the genomic sequence to be altered. In particular, these sequences are directed at the following genes: adenosine deaminase, p53, beta-globin, retinoblastoma, BRCA1, BRCA2, CFTR, cyclin-dependent kinase inhibitor 2A (CDRN2A), APC, Pactor VII, Factor IXI, haemoglobin alpha locus (HBA2), Mill, MSH2, MSH6, apolloprotein E (APOE), LDL receptor (LDLR), UDP-glucuronosyltransferase (UGII), amyloid precursor protein (APC), presentilin-1 (PSBN1) and presentilin-2 (PSBN2). These can be used in the gene therapy of diseases such as cancer, adenosine deaminase deficiency, cystic fibrosis,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The present invention provides single-stranded oligonucleotides which can
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human; gene therapy; adenosine deaminase deficiency; p53; beta-globin; retinoblastoma; BRCA1; BRCA2; CFTR; cystic fibrosis; cancer; Factor V; cyclin-dependent Kinase inhibitor 2A; CDKN2A; melanoma; APC; HBA1; HBA2; adenomatous polyposis of the colon; Factor VII; Factor IX; thrombosis; haemophilia; alpha thalassaemia; haemoglobin alpha locus 1; MLH1; APOE; mismatch repair; MSH2; MSH6; hyperlipidaemia; apolipoprotein E; LDLR; familial hypercholesterolaemia; UGT1; syndrome; APP; PSEN1; antisense; UDP-glucuronosyltransferase; amyloid precursor protein; presentiln-1; altaheimer's disease; cytostatic; antisickling; antianaemic; haemostatic; antilibemic; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Oligonucleotide for targeted alterations of genetic sequences and for treating cystic fibrosis, comprises at least one mismatch and chemical
  Alzheimer's disease, melanoma, adenomatous polyposis of the colon and various syndromes. The present sequence is one of the gene correcting oligonucleotides of the invention
                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Retinoblastoma mutation correcting oligonucleotide SEQ ID NO: 559.
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0
                                                                                                                                                 Score 9.2; DB 1; Length 17;
Pred. No. 1.6e+03;
0; Mismatches 3; Indels
                                                                                                      Sequence 17 BP; 5 A; 4 C; 2 G; 6 T; 0 U; 0 Other;
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27-MAR-2000; 2000US-0192179P.
01-JUN-2000; 2000US-026538P.
30-OCT-2000; 2000US-0244989P.
                                                                                                                                                    12.6%;
78.6%;
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                                                                                                                                                                                                                                                                                                                                                                                                                    ABA77713 standard; DNA; 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
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                                                                                                                                               Query Match
Best Local Similarity 78.6'
Matches 11, Conservative
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modification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens
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ABA77713
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New non-pathogenic HIV-1 strain carrying a deletion in its nef gene
LTR region - can be used in a vaccine to inhibit/reduce productive
infection in an individual by a pathogenic strain.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          HIV-1; AIDS; attenuation; vaccine; nef gene; avirulence; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                            HIV-1 NL4-3 nef gene nucleotide deletion 182
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12.3%; Score 9; DB 1
Local Similarity 100.0%; Pred. No. 1.3
tes 9; Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 13; Page 190; 301pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human immunodeficiency virus 1.
                                                                                                                                                                                                               BP.
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94AU-00004002.
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953 TGTATCGCTACCAA 966
                                                       4 reragcearacaaa 17
                                                                                                                                                                                                            AAQ96587 standard; DNA; 10
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schultz1-899.rng

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Hypertrophy in cardiomyocytes is treated by inhibiting function of NF-AT3. Activation of NF-AT3 mediates the calcium ion-dependent cardiac hypertrophic response to a variety of stimuli, so inhibiting it can be used to treat or prevent cardiac hypertrophy and related heart failure. Transgenic animals, or cells, containing a constitutively active NF-AT3 gene can be used as models for screening modulators of hypertrophy and for studying human disease. NF-AT3 interacts with GATA4 to have a functional role in cardiac gene expression. The BNP cardiac promoter is upregulated during cardiac hypertrophy and shows a dramatic response to the GATA4. The GATA4 were identified in the BNP promoter (SEE AAX08714-16). This
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SAGE tag; serial analysis of gene expression; antigen-presenting cell; APC; monocyte-derived dendritic cell; differential gene expression; immunostimulatory cofactor; costimulatory factor; CTL; cytotoxic T-lymphocyte; tumour antigen; immunotherapy; anticancer; ss.
                                                                                                                                                                                                                                                                                                                                               Treating hypertrophy in cardiomyocytes by inhibiting NF-A3
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                                                                                                                                                                                                                  SYSTEM.
TEXAS HEALTH SCI CENT.
                                                                                                                                                                                                                                                                        Molkentin JD;
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                                                                                                                      97US-0062864P.
97US-0065178P.
98US-0081853P.
98US-00061417.
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98US-0089844P.
98US-0089853P.
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                                                                                                                                                                                                                 UNIV TEXAS
UNIV NORTH
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15-APR-1998;
16-APR-1998;
                   WO9919471-A1
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19-JUN-1998;
19-JUN-1998;
                                                                                        15-OCT-1998;
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                                                     22-APR-1999,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Attenuation of pathogenic HIV-1 strain NL4-3 involves deletion of 1 or more decanuclectides (AAQ96406-Q97018) from the nef gene and/or 1 or more decanuclectides (AAQ96406-Q97018) from the LTR region, the sequence of AAQ96406 corresponds to nucleotides 1-10 of the nef gene (AAQ9641). The response in humans, and enable the generation of therapeutic, diagnostic and targeting agents against HIV-1 infection. (Updated on 16-OCT-2003 to standardise OS field)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NF-AT3; hypertrophy; cardiomyocytes; cardiac hypertrophic response;
heart failure; transgenic animals; screening; treatment; inhibition; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New non-pathogenic HIV-1 strain carrying a deletion in its nef gene or LTR region - can be used in a vaccine to inhibit/reduce productive infection in an individual by a pathogenic strain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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                                                                                                                                                                                                           HIV-1; AIDS; attenuation; vaccine; nef gene; avirulence; ss.
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                                                                                                                                                                          HIV-1 NL4-3 nef gene nucleotide deletion 181
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (MACF-) MACFARLANE BURNET CENT MEDICAL. (AURE-) AUSTRALIAN RED CROSS SOC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 13; Page 190; 301pp; English
                                                                                                                                                                                                                                                 Human immunodeficiency virus 1.
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94AU-00004002.
94AU-00000284.
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                                                 AAQ96586 standard; DNA; 10
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(first entry)
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Best Local Similarity
Matches 9; Conserv
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20-MAR-1996
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RESULT 25: AAX08716/

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Isolated polynucleotides differentially expressed in antigen-presenting
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                                                                                                                                 .
                                                                                                                              0; Indels
                                                        Score 9; DB 1; Length 10;
Pred. No. 1.3e+03;
Sequence 10 BP; 7 A; 1 C; 2 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human dendritic cell SAGE tag, SEQ ID NO:1326.
                                      12.3%; Scor.
100.0%; Pred. No. 1...
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They may be used in vaccines to sinduce an immune response, particularly against a tumour antigen, to modulate the genotype of an APC; to screen for agents that modulate expression of differentially expressed genes in the assessed genes, or of their encoded proteins, can be used to identify coll agnosis, prognosis and monitoring of diseases related to abnormal expression of the dendritic cell differentially expressed genes, or of their encoded proteins, can be used in active immunotherapy (or to stimulate production of a containing them are used in active immunotherapy (or to stimulate antigens and containing them are used in active immunotherapy (or to stimulate antigens and presentation of containing the encoded proteins, and secretion of cells as belonging to the monocyte lineage. Cells containing or encoded proteins, each or encoded proteins, each or encoded protei
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          9805-008991P-9805-0089991P-9805-0089991P-9805-0089991P-9805-0089997P-9805-0089997P-9805-0090035P-9805-0090043P-9805-0090044P-9805-0090044P-9805-0090044P-9805-0090044P-9805-0090044P-9805-0090044P-9805-0090044P-9805-0090044P-9805-0090044P-9805-0090044P-9805-0090044P-9805-0090044P-9805-0090044P-9805-0090044P-9805-0090044P-9805-0090044P-9805-0090044P-9805-0090044P-9805-0090044P-9805-0090047P-9805-0090044P-9805-0090047P-9805-0090047P-9805-0090044P-9805-0090047P-9805-0090047P-9805-0090047P-9805-0090047P-9805-0090047P-9805-0090047P-9805-0090047P-9805-0090047P-9805-0090047P-9805-0090077P-9805-0090077P-9805-0090077P-9805-0090077P-9805-0090077P-9805-0090077P-9805-0090077P-9805-0090077P-9805-0090077P-9805-0090077P-9805-0090077P-9805-0090077P-9805-0090077P-9805-0090077P-9805-0090077P-9805-0090077P-9805-0090077P-9805-0090077P-9805-0090077P-9805-0090077P-9805-0090077P-9805-0090077P-9805-0090077P-9805-0090077P-9805-0090077P-9805-0090077P-9805-0090077P-9805-0090077P-9805-0090077P-9805-0090077P-9805-0090077P-9805-0090077P-9805-0090077P-9805-0090077P-9805-0090077P-9805-0090077P-9805-0090077P-9805-0090077P-9805-0090077P-9805-0090077P-9805-0090077P-9805-0090077P-9805-0090077P-9805-0090077P-9805-0090077P-9805-0090077P-9805-0090077P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-009007P-9805-0090
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98US-0111715P

Roberts BL, Shankara S;

WPI; 2000-106077/09.

GENZYME CORP. ROBERTS B L. SHANKARA S.

(SHAN/) (GENZ) (ROBE/)

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Sequences AAZ77573-Z79709 represent SAGE (serial analysis of gene expression) tags used to identify mRNA transcripts encoding immunostimulatory cofactor proteins which are preferentially or differentially expressed in monocyte-derived dendritic cells compared with monocytes. Some of the transcripts correspond to known genes or ESTS (expressed sequence tags) which were previously unknown to be preferentially or differentially expressed in dendritic cells, while correspond to novel genes. Antigen-presenting cell cother transcripts correspond to novel genes. Antigen-presenting cell cother transcripts correspond to novel genes. Antigen-presenting cell cother transcribts correspond to novel genes. Antigen-presenting cell complex) and subsequent recognition by T-cell receptors is alone activation of the cytotoxic immune response, particularly against tumour cells, immunostimulatory cofactors also being required for sequences identified using the SAGE tags have several potential uses. They may be used in vaccines to induce an immune response, particularly against a tumour antigen; to modulate the genotype of an APC; to screen for against a tumour antigen; to modulate the genotype of an APC; to screen for agents that modulate expression of differential uses. They may be used in vaccines to induce an immune response, particularly against a tumour antigen; to the monocyte lineage. Cells containing these genes. Detection probes amplification primers for the copulation of antigen-specific effector cells) and vectors containing them are used in gene therapy. Co-administration of tumour antigens and presentation of co-stimulatory factors ensures adequate antigen presentation of co-stimulatory factors ensures adequate antigen generation of co-stimulatory signals, migration to femential and secretion of to election of co-centure and escretion of co-timulatory signals.

**Correction**

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cells, useful in gene vaccines against cancer.
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                             Claim 1; Page 103; 130pp; English.
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Isolated polynucleotides differentially expressed in antigen-presenting cells, useful in gene vaccines against cancer.

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expression) tags used to identify mRNA transcripts encoding expression) tags used to identify mRNA transcripts encoding immunostimulatory cofactor proteins which are preferentially or differentially expressed in monocyte-derived dendritic cells compared with monocytes. Some of the transcripts correspond to known genes or ESTS (expressed sequence tags) which were previously unknown to be preferentially or differentially expressed in dendritic cells, while preferentially or differentially expressed in dendritic cells, while other transcripts correspond to novel genes. Antigen-presenting cell other transcripts correspond to novel genes. Antigen-presenting cell other transcripts correspond to novel genes, particularly against tumour cells. Tumour antigen presentation by T-cell receptors is alone activate a robust cytocoxic immune response that can lyse the tumour cells, immunostimulatory cofactors also being required for the tumour cells, immunostimulatory cofactors also being required for efficient activation of cytotoxic T-lymphocytes (CTLS). Nucleic acid sequences identified using the SAGE tags have several potential uses. They may be used in vaccines to induce an immune response, particularly against a tumour antigen; to modulate the genotype of an APC; no screen for against that modulate expression of diseases related to abnormal expression of the dendritic cell differentially expressed genes, or of their encoded proteins, can be used in active immunotherapy (or costimulate production of a population of entigen seed in active immunotherapy (or costimulate production of a population of entigen seed in active immunotherapy (or costimulatory factors ensures adequate antigens and presentation of entigens und presentation of extimulatory factors ensures adequate antigens and presentation of costimulatory factors ensures adequate antigen presentation of extimulatory signals, migration of chemotypenses of extention of T cell growth factors and secretion of T cells generaled.
Claim 1; Page 107; 130pp; English.
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99WO-US013800, 98US-0089833P. 98US-0089844P.

18-JUN-1999; 23-DEC-1999.

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Homo sapiens. WO9965924-A2. ö

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that are preferentially transcribed in the metastatic breast tumour that are preferentially transcribed in the metastatic breast tumour cities. It is are upregalated in metastatic breast tumour cells). AAZ83942 to AAZ86677 represent tags corresponding to distinct transcripts that are preferentially transcribed in the primary or non-metastatic breast tumour cells). These tissue (i.e. are downregulated in metastatic breast tumour cells). These transcripts can be used for diagnosis, prognosis, monitoring and treatment of breast cancer, particularly where metastatic. Diagnosis is by standard immunoassays or hybridisation/amplification reactions. Compounds that modulate expression of the transcripts are potentially useful for treatment of (metastatic) breast cancer, while promoters from the transcripts are used to direct expression, in selected cell types, of c.g. therapeutic genes (also ribozymes or antisense sequences), particularly an antigen-encoding sequence for use in gene or cell-based vaccines. Polypeptides encoded by the transcripts are also useful in vaccines; (Ab). Ab are used to detect the polypeptides can be used to expand and isolate populations of educated, antigen-specific immune effecter
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                                              12.3%; Score 9; DB 1; Length 10; 100.0%; Pred. No. 1.3e+03; tive 0; Mismatches 0; Indels
            Sequence 10 BP; 1 A; 6 C; 0 G; 3 T; 0 U; 0 Other;
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(ROBE/) ROBERTS B L.
(SHAN/) SHANKARA S.
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that are preferentially transcribed in the metastatic breast tumour tissue (i.e. are upregulated in metastatic breast tumour calls). AAZ8647 represent tags corresponding to distinct transcripts that are preferentially transcribed in the primary or non-metastatic breast tumour cissue (i.e. are downregulated in metastatic breast tumour cells). These tissue (i.e. are downregulated in metastatic breast tumour cells). These transcripts can be used for diagnosis, prognosis, monitoring and transcripts can be used for diagnosis, prognosis, monitoring and transcripts can be used to their diagnosis, monitoring and transcripts can be used to their diagnosis, while promoters from the transcripts are potentially useful for treatment of (metastatic) breast cancer, while promoters from the transcripts are used to direct expression, in selected cell types, of e.g. therapeutic genes (also riboxymes or antisense sequences), particularly an antigen-encoding sequence for use in gene or cell-based contines, for diagnosing breast cancer and for raising specific antibodies (Ab). Ab are used to detect the polypeptides or as therapeutic
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cells, e.g. cytotoxic T lymphocytes, and these used for adoptive
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                                                    Sequence 10 BP; 5 A; 2 C; 2 G; 1 T; 0 U; 0 Other;
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(ROBE/) ROBERTS B L.
(SHAN/) SHANKARA S.
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agents. Host cells that produce the polypeptides can be used to expand and isolate populations of educated, antigen-specific immune effecter cells, e.g. cytotoxic T lymphocytes, and these used for adoptive immunotherapy
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(ROBE/) ROBERTS B L.
(SHAN/) SHANKARA S.
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vaccines; for diagnosing breast cancer and for raising specific antibodies (Ab). Ab are used to detect the polypeptides or as therapeutic agents. Host cells that produce the polypeptides can be used to expand and isolate populations of educated, antigen-specific immune effecter cells, e.g. cytotoxic I lymphocytes, and these used for adoptive
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SHANKARA S.
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particularly an antigen-encoding sequence for use in gene or cell-based vaccines. Polypeptides encoded by the transcripts are also useful in vaccines; for diagnosing breast cancer and for raising specific antibodies (Ab). Ab are used to detect the polypeptides or as therapeutic agents. Host cells that produce the polypeptides can be used to expand and isolate populations of educated, antigen-specific immune effecter cells, e.g. cytotoxic T lymphocytes, and these used for adoptive
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the transcripts are used to direct expression, in selected cell types, of e.g. therapeutic genes (also ribozymes or antisense sequences), particularly an antigen-encoding sequence for use in gene or cell-based vaccines. Polypeptides encoded by the transcripts are also useful in vaccines; for diagnosing breast cancer and for raising specific antibodies (Ab). Ab are used to detect the polypeptides or as therapeutic agents. Host cells that produce the polypeptides can be used to expand and isolate populations of educated, antigen-specific immune effecter cells, e.g. cytotoxic I lymphocytes, and these used for adoptive
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(SHAN/) SHANKARA S.
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Compounds that modulate expression of the transcripts are potentially useful for treatment of (metastatic) breast cancer, while promoters from the transcripts are used to direct expression, in selected cell types, of e.g. therapeutic genes (also ribozymes or antisense sequences), particularly an antigen-encoding sequence for use in gene or cell-based vaccines, Polypeptides encoded by the transcripts are also useful in autibodies (Ab). Ab are used to detect the polypeptides or as therapeutic and isolate populations of educated, antigen-specific immune effecter cells, e.g. cytotoxic T lymphocytes, and these used for adoptive immunotherapy
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treatment of breast cancer, particularly where metastatic. Diagnosis is by standard immunoassays or hybridisation/amplification reactions. Compounds that modulate expression of the transcripts are potentially useful for treatment of (metastatic) breast cancer, while promoters from the transcripts are used to direct expression, in selected cell types, of e.g. therapeutic genes (also ribozymes or antisense sequences), particularly an antigen-encoding sequence for use in gene or cell-based vaccines. Polypeptides encoded by the transcripts are also useful in vaccines; for diagnosing breast cancer and for raising specific amplibulates (Ab). Ab are used to detect the polypeptides or as therapeutic agents. Host cells that produce the polypeptides can be used to expand and isolate populations of educated, antigen-specific immune effecter cells, e.g. cytotoxic T lymphocytes, and these used for adoptive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention describes a method of identifying the type of cell in a sample, involving determining which of the sequences AAH63161-AAH64721 is expressed by the cell. The transcriptomes described in the invention are cell-type specific, cancer specific or ubiquitously expressed in humans. They can also be used to screen for drugs, reduce cancer specific gene expression, standardise expression and restore the function of a diseased cell or tissue. The present sequence is one of the transcriptomes described in the exemplification of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New isolated polynucleotides, useful for identifying specific cell type, such as cancer cell, comprises transcriptomes expressed in particular cell types.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human, transcriptome, gene expression pattern, cancer, drug screening.
cancer diagnosis, cell specific gene expression, ss.
                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human ubiquitously expressed transcriptome sequence SEQ ID NO: 644
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                                                                                                                                                                                                                                                                                                                                                12.3%; Score 9; DB 1; Length 10; 100.0%; Pred. No. 1.3e+03; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                            Sequence 10 BP; 7 A; 1 C; 2 G; 0 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAH63804 standard; cDNA; 10 BP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Vogelstein B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21-NOV-2000; 2000WO-US031922.
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                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity 100.
Matches 9; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                908 TTTTCTTTG 916
                                                                                                                                                                                                                                                                                                                                                                                                                                                     10 TTTTCTTTG 2
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                                                                                                                                                                                                                                                                          immunotherapy
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DB 1; Length

12.3%; Score 9;

Query Match

Sequence 10 BP; 4 A; 3 C; 2 G; 1 T; 0 U; 0 Other;

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Yeast gene coding sequences comprising NORF genes with serial analysis of gene expression (SAGE) tags, useful for studying, monitoring and affecting phases of the cell cycle.
                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Yeast, Saccharomyces cerevisiae; characterisation; cell cycle; NORF; nor previously assigned open reading frame; nonamnotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification; linker; PCR primer; ds.
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                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Yeast NORF gene SAGE tag oligonucleotide SEQ ID NO:879.
                        Pred. No. 1.3e+03;
Mismatches 0;
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भारत '80.001
                                                                                                                                                                                                                                                                                                              AAF34140 standard; DNA; 10 BP.
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                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Saccharomyces cerevisiae
                                                                                                         CTTTGGTCT 920
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                     Best Local Similarity
Matches 9; Conserv
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                                                                                                                                                                                                                                                    RESULT 2545
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AAC AAF3.AMX

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The present invention describes an isolated DNA molecule comprising a coding sequence of a yeast gene selected from a group of 745 NORF (not previously assigned open reading frame, or nonannotated ORF) genes comprising a SAGE (serial analysis of gene expression) tag. Also described are: (1) a method (M1) of using NORF genes to affect the cell cycle comprising administering administering a NORF gene whose expression varies by at clast 10% between any two phases of the cell cycle selected from log phase, S phase and G2/M; (2) a method (M2) for screening candidate and trugs comprising; (a) contacting a test substance with a yeast cell; and (b) monitoring expression of antifungal drug expression of antifungal gene is a candidate antifungal drug; (3) a method (M3) for identifying human genes which are involved in cell cycle progression comprising contacting human DNA with a probe which comprises at least 10 comprising contacting human DNA with a probe which comprises at least 10 comprising contacting a seast cell comprises of a NORF gene whose expression in a class of drugs having a characteristic effect on gene expression in a class of drugs having contacting a yeast cell with a candidate drug and monitoring expression in the yeast cell of at least 1 NORF gene whose expression is affected by the class of the cell with a candidate drug and monitoring expression in the yeast cell of a least 1 NORF gene whose expressed genes may be used to identify and large. The NORF genes may be used to identify and addise drugs which affect the cell cycle in methods may be used to identify andidate drugs which affect the cell cycle and for identification of antifungal drugs. AAF33268 to AAF44064 represent SAGE tags used in the exemplification of the present invention.

AAF33262 to AAF33367 represent linkers and PCR primers used in the exemplification of the present invention. Example; Page 31; 419pp; English.

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The present invention describes an isolated DNA molecule comprising a coding sequence of a yeast gene selected from a group of 745 NORF (not previously assigned open reading frame; or nonannotated ORF) genes comprising a SAGE (serial analysis of gene expression) tag. Also described are: (1) a method (M1) of using NORF genes to affect the cell cycle comprising administering a NORF gene whose expression varies by at least 10% between any two phases of the cell cycle selected from log phase, 5 phase and G2/M; (2) a method (M2) for screening candidate antifungal drugs comprising: (a) contacting a test substance which a yeast cell; and (b) monitoring expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for the yeast gene is a candidate antifungal drug; (3) a method (M3) for identifying human genes which are involved in cell cycle progression comprising contacting human DNA with a probe which comprises at least 10 comprising contacting a newtose expression varies as in M1; or identifying a candidate drug as a member of a class of drugs having a characteristic effect on gene expression in a yeast cell comprising contacting a yeast cell with a candidate drug and contacting expression in the yeast cell with a candidate drug and contacting expression in the yeast cell of at least 10 contributing expression in the yeast cell of at least 10 contributing expression in the yeast cell of at least 10 contributing expression in the yeast cell of at least 10 contributing expression of an expressed genes may be used as markers of phases of the cell cycle. The methods may be used to identify candidate drugs which affect the cell cycle and for identification of antifungal drugs. Apr31268 to AAP44064 crepresent SAGE tags used in the exemplification of the present invention.
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                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                   Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF; nor previously assigned open reading frame; nonannotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification;
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                                            Indels
    Length 10;
                                                                                                                                                                                                                                                                                                                                                                             Yeast NORF gene SAGE tag oligonucleotide SEQ ID NO:6019.
Score 9; DB 1; Le:
Pred. No. 1.3e+03;
0; Mismatches 0;
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Query Match
Best Local Similarity 100.0%; Pr
Matches 9; Conservative 0;
                                                                                                                                                                                                                                              AAF39280 standard; DNA; 10 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Saccharomyces cerevisiae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               linker; PCR primer; ds.
                                                                                         916 GGTCTTTGC 924
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                                                                                                                                    9 GGICTITGC
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The present invention describes an isolated DNA molecule comprising a coding sequence of a yeast gene selected from a group of 745 NORF (not previously assigned open reading frame, or nonamonated ORF) genes comprising a sasigned open reading frame, or nonamonated ORF) genes comprising a daministering a NORF gene expression varies by at least 10% between any two phases of the cell cycle selected from log phase, 5 phase and G2/M; (2) a method (M2) for screening candidate antifungal drugs comprising: (a) contacting a test substance with a yeast cell; and (b) monitoring expression of a NORF gene whose expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for the yeast gene is a candidate antifungal drug; (3) a method (M3) for identifying human genes which are involved in cell cycle progression contiguous nucleotides of a NORF gene whose expression varies as in M1; and (4) a method (M4) for identifying a candidate drug as a member of a class of drugs having a characteristic effect on gene expression in a cyeast cell comprising contacting a yeast cell with a candidate drug and monitoring expression in the yeast cell of at least 1 NORF gene may be used costudy, monitor and affect phases of the cell cycle, the differentially expression is affected by the class of three cell cycle, the differentially expression be used as markers of phases of the cell cycle. The
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AAF33262 to AAF33267 represent linkers and PCR primers used in the SAGE method, in the exemplification of the present invention
                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF; nor previously assigned open reading frame; nonamnotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification; linker; PCR primer; ds.
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0
                                                                                                                                    0; Indels
                                                                                            12.3%; Score 9; DB 1; Length 10; 100.0%; Pred. No. 1.3e+03; eive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                      Yeast NORF gene SAGE tag oligonucleotide SEQ ID NO:5780.
                                                          Sequence 10 BP; 1 A; 2 C; 4 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                    AAF39041 standard; DNA; 10 BP.
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                                                                             Query Match
Best Local Similarity 100..
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Saccharomyces cerevisiae.
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The present invention describes an isolated DNA molecule comprising a coding sequence of a yeast gene selected from a group of 745 NORF (not previously assigned oppen reading frame, or nonannotated ORF) genes comprising a SAGE (serial analysis of gene expression) tag. Also described are: (1) a method (M1) of using NORF gene to affect the cell cycle comprising administering a NORF gene whose expression varies by at least 10% between any two phases of the cell cycle selected from log phase, S phase and G2/M; (2) a method (M2) for screening candidate antifungal drugs comprising: (a) contacting a test substance with a yeast cell; and (b) monitoring expression of a NORF gene whose expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for identifying human pases which are in cell cycle progression contiguous nucleotides of a NORF gene whose expression varies as in M1; a method (M4) for identifying a candidate drug as a member of a class of drugs having a characteristic effect on gene expression in a yeast cell comprising concacting a yeast cell with a candidate drug and monitoring expression in the yeast cell of at least 1 NORF gene whose
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methods may be used to identify candidate drugs which affect the cell cycle and for identification of antifungal drugs. AAF33268 to AAF44064 represent SAGE tags used in the exemplification of the present invention. AAF33262 to AAF33267 represent linkers and PCR primers used in the SAGE method, in the exemplification of the present invention
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                                                                                                                    Sequence 10 BP; 5 A; 1 C; 3 G; 1 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                          AAF36893 standard; DNA; 10 BP.
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AAF36893/c
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The present invention describes an isolated DNA molecule comprising a coding sequence of a yeast gene selected from a group of 745 NORF (not previously assigned open reading frame, or nonanotated ORF) genes comprising a sasigned open reading frame, or nonanotated ORF) genes comprising a sasigned open reading frame, or nonanotated ORF) genes comprising administering a NORF gene expression varies by at least 10% between any two phases of the cell cycle selected from log phase, S phase and G2/M; (2) a method (M2) for screening candidate or infilmad drugs comprising: (a) contacting a test substance with a yeast cell; and (b) monitoring expression of a NORF gene whose expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for identifying human genes which are involved in cell cycle progression contiguous mucleotides of a NORF gene whose expression contiguous mucleotides of a NORF gene which comprises at least 10 contiguous mucleotides of a NORF gene whose expression varies as in M1; and (4) a method (M4) for identifying a candidate drug as a member of a
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expression is affected by the class of drugs. The NORF genes may be used to study, monitor and affect phases of the cell cycle, the differentially expressed genes may be used as markers of phases of the cell cycle. The methods may be used to identify candidate drugs which affect the cell cycle and for identification of antifungal drugs. AAP33268 to AAF4064 represent inhere samplification of the present invention. PAF33262 to AAF33267 represent linkers and PCR primers used in the SAGE method, in the exemplification of the present invention.
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                                                                                                                                                                                                                                          Query Match 12.3%; Score 9; DB 1; Length 10; Best Local Similarity 100.0%; Pred. No. 1.3e+03; Matches 9; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                     Sequence 10 BP; 5 A; 2 C; 2 G; 1 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAF42052 standard; DNA; 10 BP.
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                                                                                                                                                                                                                                                                                                                                   903 GGTCATTTT 911
                                                                                                                                                                                                                                                                                                                                                                               GGTCATTTT 1
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class of drugs having a characteristic effect on gene expression in a monitoring expression in the yeast cell of at least 1 NORF gene whose expression is affected by the class of drugs. The NORF genes may be used to study, monitor and affect phases of the cell cycle, the differentially expressed genes may be used as markers of phases of the cell cycle. The methods may be used to identify candidate drugs which affect the cell cycle and for identification of antifungal drugs. AAF33268 to AAF44064 represent inhers and PCR primers used in the exemplification of the present invention. AAF33262 to AAF33267 represent linkers and PCR primers used in the SAGE method, in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present invention describes an isolated DNA molecule comprising a coding sequence of a yeast gene selected from a group of 745 NORF (not previously assigned open reading frame, or nonamnotated ORF) genes comprising a SAGE (serial analysis of gene expression) tag. Also described are: (1) a method (M1) of using NORF genes to affect the cell cycle comprising administering a NORF gene whose expression varies by at chast 10% between any two phases of the cell cycle selected from log phase, S phase and G2/M; (2) a method (M2) for screening candidate antifungal drugs comprising: (a) contacting a test substance with a yeast cell; and (b) monitoring expression of a NORF gene whose expression varies as in M1, where a test substance which modifies the expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for identifying human genes which are involved in cell cycle progression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Yeast gene coding sequences comprising NORF genes with serial analysis of gene expression (SAGE) tags, useful for studying, monitoring and affecting phases of the cell cycle.
                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Yeast, Saccharomyces cerevisiae; characterisation; cell cycle; NORF; nor previously assigned open reading frame; nonannotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification;
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0
                                                                                                                                                                                                                                                                                                   12.3%; Score 9; DB 1; Length 10; 100.0%; Pred. No. 1.3e+03; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Yeast NORF gene SAGE tag oligonucleotide SEQ ID NO:7150.
                                                                                                                                                                                                                                                              Sequence 10 BP; 4 A; 1 C; 4 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Kinzler K;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Example, Page 255; 419pp, English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAF40411 standard; DNA; 10 BP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Velculescu V, Vogelstein B,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      14-JUN-2000; 2000WO-US016223
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  99US-00335032
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
                                                                                                                                                                                                                                                                                                                                                9; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   linker; PCR primer; ds.
                                                                                                                                                                                                                                                                                                                                                                                           928 TTATCCCTC 936
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                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
Matches 9; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAF40411;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 2
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computating contacting human DNA with a probe which comprises at least 10 contiguous nuclectides of a NORF gene whose expression varies as in M1; and (4) a method (M4) for identifying a candidate drug as a member of a class of drugs having a characteristic effect on gene expression in a monitoring expression in the yeast cell of at least 1 NORF gene whose expression is affected by the class of drugs. The NORF genes may be used to study, monitor and affect phases of the cell cycle, the differentially methods may be used to identify candidate drugs which affect the cell cycle and for identify candidate drugs which affect the cell cycle and for identify candidate drugs which affect the cell cycle and for identify candidate drugs. AAP33268 to AAF44664 represent inthe exemplification of the present invention. AAF33267 represent linkers and PCR primers used in the SAGE method, in the exemplification of the present invention.

Sequence 10 BP; 1 A; 1 C; 0 G; 8 T; 0 U; 0 Other;

12.3%; Score 9; DB 1; Length 10; 100.0%; Pred. No. 1.3e+03; trive 0; Mismatches 0; Indels Conservative 907 ATTITITIT 915 1 ATTITCTT 9 Local Similarity hes 9; Conserv Query Match Matches à

Gaps . 0

0; Indels

RESULT 2551

AAF40134 standard; DNA; 10 BP. AAF40134;

(first entry) 23-MAR-2001

Yeast NORF gene SAGE tag oligonuclectide SEQ ID NO:6873.

Yeast; Saccharomyces cerevisiae; characterisation; cell cycle; NORF; nor previously assigned open reading frame; nonannotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification; linker; PCR primer; ds.

Saccharomyces cerevisiae.

WO200077214-A2.

21-DEC-2000.

14-JUN-2000; 2000WO-US016223

99US-00335032 16-JUN-1999;

SNINGO NIND (OCYU)

Velculescu V, Vogelstein B,

WPI; 2001-061874/07.

Kinzler

Yeast gene coding sequences comprising NORF genes with serial analysis gene expression (SAGE) tags, useful for studying, monitoring and affecting phases of the cell cycle.

Example; Page 245; 419pp; English.

The present invention describes an isolated DNA molecule comprising a coding sequence of a yeast gene selected from a group of 745 NORF (not previously assigned open reading frame; or nonannotated ORF) gene comprising a SAGE (serial analysis of gene expression) tag. Also escribed are: (1) a method (M1) of using NORF genes to affect the cell cycle comprising administering a NORF gene whose expression varies by at least 10% between any two phases of the cell cycle selected from log analyses, S phase and G2/M; (2) a method (M2) for screening and and G2/M; (2) a method (M2) for screening and candidate antifungal and d2/M; (3) contacting a test substance with a yeast cell; and (b) monitoring expression of a NORF gene whose expression

the yeast gene is a candidate antifungal drug; (3) a method (M3) for identifying human genes which are involved in cell cycle progression comprising contacting human para inth a probe which comprises at least in comprising contacting human DNA with a probe which comprises at least in contiguous nucleotides of a NORF gene whose expression varies as in M1; and (4) a method (M4) for identifying a candidate drug as a member of a class of drugs having a characteristic effect on gene expression in a yeast cell comprising contacting a yeast cell with a candidate drug and monitoring expression in the yeast cell of at least 1 NORF gene whose expression is affected by the class of the cell cycle, the differentially expressed genes may be used as markers of phases of the cell cycle. The methods may be used to identify candidate drugs which affect the cell cycle and for identification of antifungal drugs. MAF33268 to AAF44664 represent SAGE tags used in the exemplification of the present invention.

ARF33262 to AAF33267 represent linkers and PCR primers used in the SAGE method, in the exemplification of the present invention.

Sequence 10 BP; 3 A; 3 C; 3 G; 1 T; 0 U; 0 Other;

Gaps ; 12.3%; Score 9; DB 1; Length 10; 100.0%; Pred. No. 1.3e+03; tive 0; Mismatches 0; Indels 9; Conservative Query Match Best Local Similarity Matches

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ઠ g RESULT 2552 AAF41681

AAF41681 standard; DNA; 10 BP

(first entry) 23-MAR-2001

AAF41681;

Yeast NORF gene SAGE tag oligonucleotide SEQ ID NO:8420.

Yeast, Saccharomyces cerevisiae; characterisation; cell cycle; NORF; nor previously assigned open reading frame; nonannocated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification; linker; PCR primer; ds.

Saccharomyces cerevisiae.

WO200077214-A2.

21-DEC-2000.

14-JUN-2000; 2000WO-US016223.

99US-00335032 16-JUN-1999;

(UYJO) UNIV JOHNS HOPKINS.

Kinzler K; Vogelstein B, Velculescu V,

WPI; 2001-061874/07.

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Yeast gene coding sequences comprising NORF genes with serial analysis of gene expression (SAGE) tags, useful for studying, monitoring and affecting phases of the cell cycle.

Example; Page 300; 419pp; English.

The present invention describes an isolated DNA molecule comprising a coding sequence of a yeast gene selected from a group of 745 NORF (not previously assigned open reading frame; or nonannotated ORF) genes comprising a SAGE (serial analysis of gene expression) tag. Also described are: (1) a method (M1) of using NORF genes to affect the cell cycle comprising administering a NORF gene whose expression varies by at least 10% between any two phases of the cell cycle selected from log

phase, S phase and G2/W; (2) a method (M2) for screening candidate antifungal drugs comprising: (a) contacting a test substance with a yeast cell; and (b) monitoring expression of a NORF gene whose expression craries as in M1, where a test substance which modifies the expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for identifying human genes which are involved in cell cycle progression of comprising contacting human DNA with a probe which comprises at least 10 contiguous nucleotides of a NORF gene whose expression varies as in M1; and (4) a method (M4) for identifying a candidate drug as a member of a class of drugs having a characteristic effect on gene expression in a yeast cell comprising contacting a yeast cell with a candidate drug and monitoring expression in the yeast cell of at least 1 NORF genes whose compression in the yeast cell of at least 1 NORF genes may be used to dentify candidate drugs which affect the cell cycle and for identification of antifungal drugs. AAF31368 to AAF44064 represent SAGE test as a north of the present invention. AAF33262 to AAF33267 represent linkers and PCR primers of method, in the exemplification of the present invention Sequence 10 BP; 5 A; 1 C; 3 G; 1 T; 0 U; 0 Other; \$

ö 0; Indels 12.3%; Score 9; DB 1; Length 10; 100.0%; Pred. No. 1.3e+03; cive 0; Mismatches 0; Indels Query Match
Best Local Similarity 100...
5. Conservative

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904 GTCATTTTC 912 9 GTCATTTTC 1 ઠે 셤

AAF40571 standard; DNA; 10 BP.

AAF40571;

23-MAR-2001 (first entry)

Yeast NORF gene SAGE tag oligonucleotide SEQ ID NO:7310.

Yeast, Saccharomyces cerevisiae; characterisation, cell cycle, NORF, nor previously assigned open reading frame; nonannotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification; linker; PCR primer; ds.

Saccharomyces cerevisiae.

WO200077214-A2

21-DEC-2000,

14-JUN-2000; 2000WO-US016223.

99US-00335032 16-JUN-1999;

SNINGO UNIV (OLYU)

Kinzler K; Velculescu V, Vogelstein B,

WPI; 2001-061874/07.

Yeast gene coding sequences comprising NORF genes with serial analysis of gene expression (SAGE) tags, useful for studying, monitoring and affecting phases of the cell cycle.

Example; Page 261; 419pp; English.

Yeast gene coding sequences comprising NORF genes with serial analysis of gene expression (SAGE) tags, useful for studying, monitoring and affecting phases of the cell cycle.

Velculescu V, Vogelstein B, Kinzler K;

WPI; 2001-061874/07.

UYJO) UNIV JOHNS HOPKINS

16-JUN-1999;

The present invention describes an isolated DNA molecule comprising

Example; Page 244; 419pp; English.

The present invention describes an isolated DNA molecule comprising a coding sequence of a yeast gene selected from a group of 745 NORF (not previously assigned open reading frame; or nonamotated ORF) genes comprising a SAGE (serial analysis of gene expression) tag. Also

described are: (1) a method (M1) of using NORF genes to affect the cell cycle comprising administering a NORF gene whose expression varies by at least 10% between any two phases of the cell cycle selected from log phase, S phase and G2/M; (2) a method (M2) for screening candidate antifungal drugs comprising: (a) contacting a test substance with a yeast cell; and (b) monitoring expression of varies as in M1, where a test substance which modifies the expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for identifying human genes which a probe which comprises at least 10 comprising contacting human DNA with a probe which comprises at least 10 contiguous nucleotides of a NORF gene whose expression varies as in M1; and (4) a method (M4) for identifying a candidate drug as a member of contiguous nucleotides of a NORF gene whose expression in a class of drugs having a characteristic effect on gene expression in a class of drugs having a characteristic effect on gene expression in a class of drugs having a class of drugs. The NORF gene whose expression is affected by the class of drugs. The NORF gene whose cut dry, monitor and affect phases of the cell cycle. The methods may be used to identify candidate drugs which affect the cell cycle and for identification of antifungal drugs. Apfalled the represent invention. ö Gaps Yeast, Saccharomyces cerevisiae; characterisation; cell cycle; NORF; nor previously assigned open reading frame; nonannotated ORF; SAGE; serial analysis of gene expression; antifungal; tag; identification; . 0 12.3%; Score 9; DB 1; Length 10; 100.0%; Pred. No. 1.3e+03; ative 0; Mismatches 0; Indels Yeast NORF gene SAGE tag oligonucleotide SEQ ID NO:6858. method, in the exemplification of the present invention Sequence 10 BP; 6 A; 3 C; 1 G; 0 T; 0 U; 0 Other; AAF40119 standard; DNA; 10 BP. 14-JUN-2000; 2000WO-US016223. 23-MAR-2001 (first entry) Conservative Saccharomyces cerevisiae. linker; PCR primer; ds. 918 ~ Query Match Best Local Similarity Matches 9; Conserv 910 TTCTTTGGT 10 TICTITGGI WO200077214-A2. 21-DEC-2000. AAF40119; RESULT 2554 AAF40119 d

Example, Page 182; 419pp; English.

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coding sequence of a yeast gene selected from a group of 745 NORF (not previously assigned open reading frame, or nonannotated ORF) genes comprising a SAGE (serial analysis of gene expression) rag, Also comprising administering a NORF gene whose expression varies by at least 10% between any two phases of the cell cycle selected from log phase, S phase and G2/M; (2) a method (M2) for screening candidate by at call; and (D) monitoring expression of a NORF gene whose expression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for identifying human genes which are involved in cell cycle progression of the yeast gene is a candidate antifungal drug; (3) a method (M3) for identifying human DNA with a probe which comprises at least 10 contiguous nucleotides of a NORF gene whose expression varies as in M1; an ethod (M4) for identifying a candidate drug as a member of contiguous nucleotides of a NORF gene whose expression in a yeast cell comprising contacting a yeast cell with a candidate drug as member of a yeast cell comprising contacting a yeast cell with a candidate drug and monitoring expression in the yeast cell of at least 1 NORF gene whose cyclession is affected by the class of drugs. The NORF genes may be used to identify candidate drugs which affect the cell cycle. The methods may be used to identify candidate drugs which affect the cell cycle and for identification of antifungal drugs. AAF33568 to AAF44064 represent SAGE tags used in the exemplification of the present invention. AAF33262 to AAF33262 to AAF33262 to AAF33262 to AAF33262 to AAF33262 to AAF33264 to AAF44064 represent invention of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         method, in the exemplification of the present invention
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Sequence 10 BP; 1 A; 3 C; 1 G; 5 T; 0 U; 0 Other;

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12.3%; Score 9; DB 1; Length 10; 100.0%; Pred. No. 1.3e+03; ive 0; Mismatches 0; Indels
Query Match
12.3
Best Local Similarity 100.
Matches 9; Conservative
                                                                               918 TCTTTGCCT 926
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Gaps

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10 rciridecr

RESULT 2555 AAF38371

AAF38371;

AAF38371 standard; DNA; 10 BP.

(first entry) 23-MAR-2001 Yeast, Saccharomyces cerevisiae, characterisation, cell cycle, NORF, nor previously assigned open reading frame; nonannotated ORF, SAGE, serial analysis of gene expression; antifungal; tag; identification;

Yeast NORF gene SAGE tag oligonucleotide SEQ ID NO:5110.

Saccharomyces cerevisiae

linker; PCR primer; ds.

WO200077214-A2

21-DEC-2000.

14-JUN-2000; 2000WO-US016223

99US-00335032 16-JUN-1999;

UNIO) UNIV JOHNS HOPKINS

Kinzler K; Vogelstein B, Velculescu V,

WPI; 2001-061874/07.

Yeast gene coding sequences comprising NORF genes with serial analysis of gene expression (SAGE) tags, useful for studying, monitoring and affecting phases of the cell cycle.

```
The present invention describes an isolated DNA molecule comprising a coding sequence of a yeast gene selected from a group of 745 NORF (not perviously assigned open reading frame; or nonannotated ORF) genes comprising a SAGE (serial analysis of gene expression) tag. Also described are: (1) a method (M1) of using NORF genes to affect the cell cycle comprising administering a NORF gene whose expression varies by at contacting a test substance with a yeast castifulated comprising; (a) contacting a test substance with a yeast call; and (b) monitoring expression of a NORF gene whose expression of the yeast gene is a candidate antifungal drug; (3) a method (M2) for cell; and (b) monitoring expression of a NORF gene whose expression of varies as in M1, where a test substance which modifies the expression of centrifying human genes which are involved in cell cycle progression of identifying human genes which are involved in cell cycle progression of contiguous nucleotides of a NORF gene whose expression varies as in M1, and (4) a method (M4) for identifying a candidate drug as a member of a class of drugs having a characteristic effect on gene expression in a contiguous nucleotides of a NORF gene whose expression is affected by the class of the cell of at least 1 NORF genes may be used controlly monitoring expression in the yeast cell with a candidate drug and monitoring expression in the yeast cell of at least 1 NORF genes may be used to identify candidate drugs which affect the cell cycle and for identification of antifungal drugs. The NORF genes may be used to identify candidate drugs which affect the cell cycle and for identification of antifungal drugs. The NORF genes in the exemplification of the present invention.

AAF31262 to AAF31367 represent linkers and por immers used in the seemplification of the present invention.
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; 0 0; Indels 12.3%; Score 9; DB 1; Length 10; 100.0%; Pred. No. 1.3e+03; ive 0; Mismatches 0; Indels Sequence 10 BP; 0 A; 1 C; 3 G; 6 T; 0 U; 0 Other; Query Match Best Local Similarity 100. Matches 9, Conservative

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Gaps

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RESULT 2556

AAF36038 standard; DNA; 10 BP. AAF36038; Yeast NORF gene SAGE tag oligonucleotide SEQ ID NO:2777.

23-MAR-2001 (first entry)

Yeast, Saccharomyces cerevisiae, characterisation, cell cycle, NORF, nor previously assigned open reading frame, nonannotated ORF, SAGE, serial analysis of gene expression, antifungal, tag, identification, linker, PCR primer, ds.

Saccharomyces cerevisiae

WO200077214-A2.

21-DEC-2000

14-JUN-2000; 2000WO-US016223.

16-JUN-1999;

(UYJO) UNIV JOHNS HOPKINS.

Kinzler K; Jelculescu V, Vogelstein B,

WPI; 2001-061874/07.

Yeast gene coding sequences comprising NORF genes with serial analysis of

gene expression (SAGE) tags, useful for studying, monitoring and affecting phases of the cell cycle.

The present invention describes an isolated DNA molecule comprising a coding sequence of a yeast gene selected from a group of 745 NORF (not previously assigned open reading frame, or nonanotated ORF) genes comprising a SAGE (serial analysis of gene expression) tag. Also described are: (1) a method (M1) of using NORF genes whose expression varies by at cycle comprising administering an NORF gene whose expression varies by at antifungal drugs comprising; (a) amethod (M2) for screening candidate antifungal drugs comprising; (a) contacting a test substance with a yeast call; and (b) monitoring expression of a test substance which modifies the expression of the yeast gene is a candidate antifungal drug; (3) amethod (M3) for identifying human genes which are involved in cell cycle progression comprising contacting human DNA with a probe which comprises at least 10 and (4) a method (M4) for identifying a candidate drug as a member of a class of drugs having a characteristic effect on gene expression in a class of drugs having ocharacting a gene candidate drug as a member of a class of drugs and fect phases of the cell cycle. The methods may be used to study, monitor and affect phases of the cell cycle of study, monitor and affect phases of the cell cycle and for identification of antifungal drugs. Affect the cell cycle and for identification of antifungal drugs, which affect the cell cycle and for identification of antifungal drugs. Affect the cell cycle and for identification of antifungal drugs, which affect the cell cycle and for identification of antifungal drugs. Affect the cell cycle and for identification of antifungal drugs. Affect the creamplance of the present invention. Affect to AFF44064

The expression in the exemplification of the present invention. AFF3262 to AFF46064 ö Gaps o; 0; Indels 12.3%; Score 9; DB 1; Length 10; 100.0%; Pred. No. 1.3e+03; Live 0; Mismatches 0; Indels Sequence 10 BP; 6 A; 2 C; 1 G; 1 T; 0 U; 0 Other; Example; Page 99; 419pp; English 9; Conservative Query Match Best Local Similarity Best Loca Matches

913 TTTGGTCTT 921 TTTGGTCTT 10 ઠ g

ABK68693 standard; DNA; 10 BP. 2557 ABKG 8693
ABKG 8693
ABKG 8693
AC ABKG

(first entry) 02-JUL-2002 ABK68693;

Human SCYA2 gene allele-specific oligonucleotide PCR primer #1.

Human, small inducible cytokine A2; SCYA2; primer; ss; haplotype pair; haplotyping; atherosclerosis; antiarteriosclerotic; gene therapy; single nucleotide polymorphism; genotyping; drug screening; PCR; chromosome 17q11.2-q21.1.

Homo sapiens

WO200218413-A2

07-MAR-2002.

28-AUG-2001; 2001WO-US026899.

28-AUG-2000; 2000US-0228496P

(GENA-) GENAISSANCE PHARM INC.

Ħ Lee Kumar AM, Finkel K, Koshy B, Anastasio AE,

The invention relates to single nucleotide polymorphisms in the gene encoding human small inducible cytokine A2 (SCYA2) polypeptide. A method conding human small inducible cytokine A2 (SCYA2) polypeptide. A method conclude at one or more polymorphic sites and determining whether one of the specification or whether both copies are defined by a haplotypes given the specification or whether both copies are defined by a haplotype pair. This method is useful in genotyping, whereby all possible haplotype pair. This method is useful in genotyping, whereby all possible haplotype can be trained by comparing the frequency of the haplotype or haplotype or haplotype pair of the SCYA2 gene can be trained by comparing the frequency of the haplotype or haplotype pair in a reference population, where a higher haplotype or haplotype pair in a reference population, where a higher haplotype or haplotype pair. SCYA2 and its corresponding DNA are used for studying the expression and function of SCYA2, and in screening for candidate drugs to treat diseases related to SCYA2 activity, such as a therosclerosis. Sequences ABX68631-ABX6804 represent allele-specific oligonucleotide PCR primers used for detecting SCYA2 gene polymorphisms New genetic variants having polymorphisms in the small inducible cytokine Al (SCYA2) gene, useful for studying the function of SCYA2 , and for treating disorders affected by expression or function of the SCYA2 Gaps ö 12.3%; Score 9; DB 1; Length 10; llarity 100.0%; Pred. No. 1.3e+03; Conservative 0; Mismatches 0; Indels Sequence 10 BP; 0 A; 7 C; 0 G; 3 T; 0 U; 0 Other; Claim 19; Page 13; 58pp; English. 939 Query Match Best Local Similarity Matches 9; Conserv 931 TCCCTCCTC

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1034/c ABL99034 standard; cDNA; 10 BP. RESULT 2558

ABL99034;

25-JUN-2002 (first entry)

Mouse neuronal regeneration related SAGE EST 29

Mouse; neuronal; regeneration; nerve cell; synaptic efficiency; memory; learning disorder; serial analysis of gene expression; SAGE; gene expression; hippocampus; expressed sequence tag; EST; ss.

Mus sp.

DE10048893-A1.

11-APR-2002.

02-OCT-2000; 2000DE-01048893

02-OCT-2000; 2000DE-01048893

(LION-) LION BIOSCIENCE

WPI; 2002-341428/38.

New nucleic acids involved in neuronal regeneration, useful in screening for modulators of regeneration or synaptic efficiency, and potential therapeutic agents.

Example 6; Page 9; 38pp; German.

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The invention describes an isolated human pyridoxal (pyridoxine, vitamin B6) kinase, (PDXK) polynuclectide. The polynuclectide is useful in studying the expression and function of PDXK, and in expressing PDXK protein for use in screening for candidate drugs to treat PDXK calated diseases and for therapeutic purposes. A transgenic animal is useful for studying expression of the PDXK isogenes in vivo, for in vivo screening and testing of drugs targeted against PDXK protein, and for testing the efficacy of therapeutic agents and compounds for autoimmune polyglandular disease type 1. The polypeptide is useful for studying the effect of the variation on the biological activity of PDXK and the binding affinity of candidate drugs targeting PDXK for the treatment of autoimmune polyglandular disease type 1. Genotyping and haplotyping is useful for improving the efficacy and reliability of several steps in the discovery and development of drugs for treating diseases associated with PDXK activity, e.g., autoimmune polyglandular disease type 1, to validate PDXK
                                                                                                                                                                                                                                                                                                                          ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Isolated human pyridoxal (pyridoxine, vitamin B6) kinase polyNts, useful for therapeutic purposes, for studying the expression and function of the polyNt, and for expressing pyridoxal protein.
The invention relates to nucleic acids (ABL98957-ABL99004) involved in regenerative neuronal processes and encoded proteins (ABB79405-ABB79409) used to screen for compounds and potential therapeutic agents that modulate nerve cell regeneration and/or synaptic efficiency. They may also be used for treatment or diagnosis of defective or pathological memory and learning conditions. The present sequence is that of an isolated from serial analysis of gene expression (SAGE) experiments comparing gene expression in the hippocampus of GFAP/L1 transgenic mice versus a wildtype control. The resultant EST were used to isolate the nucleic acids of the invention
                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Pyridoxal kinase; pyridoxine; vitamin B6;
PDXK autoimmune polyglandular disease type 1; transgenic animal;
gene therapy; primer extension; primer; ss.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pyridoxal (Pyridoxine, vitamin B6) Kinase (PDXK) primer #13
                                                                                                                                                                                                                                                                           12.3%; Score 9; DB 1; Length 10; 100.0%; Pred. No. 1.3e+03; cive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                      0; Indels
                                                                                                                                                                                                                                        Sequence 10 BP; 5 A; 3 C; 1 G; 1 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABK16990 standard; DNA; 10 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               24-MAY-2001; 2001WO-US016909.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24-MAY-2000; 2000US-0206664P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (first entry)
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                                                                                                                                                                                                                                                                                                                      9; Conservative
                                                                                                                                                                                                                                                                                                                                                                940 TICATIGGT 948
                                                                                                                                                                                                                                                                                                                                                                                                      9 TTCATTGGT 1
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                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO200190125-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    26-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         29-NOV-2001.
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                                                                                                                                                                                                                                                                                                                   Matches
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ABK16990/
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The invention relates to SAGE (serial analysis of gene expression) tags
correpresenting groups of genes which are differentially expressed in human
correpresenting groups of genes which are differentially expressed in human
correctlular carcinoma (HCC) compared with normal human liver tissue.
Cocated downstream of the 5' CATG-3' sequence motif lying nearest to the
colour downstream of the 5' CATG-3' sequence motif lying nearest to the
colour downstream of the 5' CATG-3' sequence motif lying nearest to the
colour of cDNAs derived from a variety of genes. These tags serve
counquely identify each transcript and can thus be used to analyse the
colour of cDNAs derived from a variety of genes. These tags serve
control of cDNAs derived from a variety of genes. These tags serve
control of cDNAs derived from a variety of genes. The invention also
control of cDNAs derived from a variety of genes. The invention also
control of proups of genes expressed in chronic hepatitis C
control of groups of genes that are overexpressed in chronic
chaptitis C liver tissue or HCC. Groups of genes differentially expressed
control of drugs to treat chronic hepatitis C and/or HCC. Sequences
controlly, and antibodies against the gene products may be used in the
control of drugs to treat chronic hepatitis C and/or HCC. Sequences
controlly and antibodies against the gene products may be used
controlly abbused genes out of those genes which are overexpressed in chronic
controlly expressed genes which are overexpressed in chronic
controlly expressed dense which are overexpressed in chronic
                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SAGE tag; serial analysis of gene expression; human; chronic hepatitis C; CH; liver tissue; hepatocellular carcinoma; cancer; tumour; HCC; expression pattern; differential expression; ss.
                                                                      β
as a candidate agent for treating a specific condition or disease predicted to be associated with PDXK activity, and in the design of clinical trials of candidate drugs. This sequence is one of 38 (see ABK16978-ABK17015) primers used for detecting PDXK gene polymorphisms primer extension terminates, described in the method of the invention
                                                                                                                                                                                                                                    Gaps
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                                                                                                                                                                                    12.3%; Score 9; DB 1; Length 10; 100.0%; Pred. No. 1.3e+03; tive 0; Mismatches 0; Indels
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                                                                                                                                          Sequence 10 BP; 8 A; 1 C; 1 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BP; 5 A; 2 C; 2 G; 1 T; 0 U; 0 Other,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (KAGA-) KAGAKU GLJUTSU SHINKO JIGYODAN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; Page 10; 139pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                            ABV84222 standard; cDNA; 10 BP.
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                                                                                                                                                                                                         Local Similarity 100.
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Matches
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The present invention relates to novel single nucleotide polymorphisms (SNPs) in the human bone morphogenetic protein receptor type II (SNPs) in the human bone morphogenetic protein receptor type II and methods for haploryping and/or genotyping the BMPR2 gene. The methods of the invention make use of allele-specific oligonucleotides (ASOS) as probes and primers, and/or primer-extension oligonucleotides for detecting the BMPR2 gene polymorphisms. The polymucleotides for compounds are useful for the treatment of diseases associated with BMPR2 activity, such as primary pulmonary hypertension (PPH) and bone disorders, ABK54467-ABK5482 represent primer-extension oligonucleotides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel isolated polynucleotide which is a polymorphic variant of bone morphogenetic protein receptor, type II (serine/threonine kinase) (BMPR2) gene useful for expressing BMPR2 protein isoform used in drug screening.
                                                        Primer-extension oligonucleotide #6 to detect human BMPR2 polymorphisms
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human, enolase 3(beta, muscle); ENO3; single nucleotide polymorphism;
SNP; haplotype analysis; isogene; primer; ss.
                                                                                                 Human, single nucleotide polymorphism, SNP, BMPR2; chromosome 2q33-ebone morphogenetic protein receptor type II; serine/threonine kinas haplotyping; genetyping; gene; primary pulmonary hypertension; PPH; bone disorder; primar; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human enolase 3 gene allele specific primer SEQ ID NO: 58.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               for detecting human BMPR2 gene polymorphisms
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sanchis A;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 18; Page 15; 98pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                              (GENA-) GENAISSANCE PHARM INC. (LANZ/) LANZ E M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Messer C,
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                                                                                                                                                                                                                                                                                                                                            27-AUG-2001; 2001WO-US026641.
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                      05-JUN-2002 (first entry)
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Matches 9; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention relates to a nucleic acid library comprising mycdependent downstream genes or their functional fragments essentially capable of supporting a neoplastic character of cancer such as growth, invasion or spread. These myc target or tag sequences are identified by SAGE (serial analysis of gene expression). The library is useful to new diagnoses and treatments for cancer. The invention is also useful to enhance production of recombinant proteins in a production system with high expression of endogenous or transfected myc oncogenes. ABK21412-ABK23828 represent transcript tag DNA sequences that are activated or repressed by N-myc in human neuroblastoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A new nucleic acid library of myc-dependent downstream genes capable of supporting a neoplastic characteristic of cancer is useful to find new therapies and diagnoses for cancer.
                                                                                                                                                                                                                                                                                                                                                                                                          Myc-dependent downstream gene; neoplastic; cancer; growth; invasion; spread; myc target; myc tag; SAGE; serial analysis of gene expression; myc oncogene; N-myc; human neuroblastoma; cytostatic; ds.
                                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                     Transcript tag DNA sequence #199 induced or suppressed by N-myc.
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  Length 10;
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12.3%; Score 9; DB 1; Let 100.0%; Pred. No. 1.3e+03; ive 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Disclosure; Page 54; 69pp; English
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29-JUN-2000; 2000EP-00202284.
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                                                                                                                                                                                                                                        ABK23610 standard; DNA; 10
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  Query Match 12.3
Best Local Similarity 100.
Matches 9; Conservative
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                                                                                                                               9 CTTTGGTCT 1
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RESULT 2562 ABKS4472 ID ABK54473 XX AC ABKS4473

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ADE13989 standard; DNA; 10
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                                                                   Local Similarity 100.
nes 9; Conservative
                                                                                                 931 TCCCTCCTC 939
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                                                                                                                    TCCCTCCTC
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RAYMOND V.
                                                                                                                                                                                                                                                                                                            US2003190617-A1
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                                                           Query Match
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                                                                                                                                                                            The present invention provides the protein, cDNA and genomic sequences of a human enclase 3 (beta, muscle) isogene containing a number of single nucleotide polymorphisms (SNPs). The sequences can be used to identify the haplotype of an individual and identify whether particular haplotypes are linked to certain diseases. The present sequence is a primer for the ENO3 gene described in the exemplification of the invention
                                                                                                                  Novel genetic variants of enolase 3, (beta, muscle) gene useful in studying expression and function of the protein, and for screening drugs to treat disorders of glycolytic pathway.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The invention relates to novel monobactam compounds. A compound of the invention has antibacterial activity, and acts as a PBP2a inhibitor. The compounds are used as antibacterial agents. The monobactam compounds
                                                                                                                                                                                                                                                                                                                                                                                                                                                             ss; monobactam; antibacterial; PBP2a; inhibitor; methicillin resistant Staphylococcus aureus; MRSA; lactam antibiotic.
                                                                                                                                                                                                                                                                                          Gaps
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                                                                                                                                                                                                                                                                     12.3%; Score 9; DB 1; Ler 100.0%; Pred. No. 1.3e+03;
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                                                                                                                                                                                                                                                                                          Mismatches
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                                                                               Parks KE;
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                                                                                                                                                         Claim 18; Page 14; 90pp; English.
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ADC17774 standard; DNA; 10 BP.
                                                          (GENA-) GENAISSANCE PHARM INC
                                                                              Koshy B,
                   02-JUL-2001; 2001WO-US020952.
                                      30-JUN-2000; 2000US-0215236P.
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                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
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Best Local Similarity 100.
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                                                                                                 WPI; 2002-154721/20.
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                                                                             Finkel K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO2003051314-A2.
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10-JAN-2002
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The invention relates to an isolated nucleic acid (NI) comprising at least 20 but not more than 1500 consecutive nucleotides of the optineurin promoter appearing as ADBI3800. Also included are the optineurin promoter operably linked to a heterologous nucleic acid, a nucleic acid capable of detecting a single nucleotide polymorphism (SNP) in the optineurin promoter, a host cell comprising the promoter operably linked to a heterologous sequence, diagnosing or prognosing glaucoma in a sample obtained from a cell or bodily fluid (comprising glaucoma in a sample obtained from a cell or bodily fluid (comprising detecting a polymorphism in a promoter region of the optineurin gene, associated with a glaucoma in a promoter region of the optineurin promoter sequence variation in a sample containing a SNP sequence variation in a sample containing DNA, detecting the presence of an optineurin promoter sequence variation in a sample containing in loss of visual field in a patient (or the severity or progression of glaucoma in a patient, comprising providing amplification reaction primers that direct amplification of a selected amplification reaction primers that direct amplification of a selected concleir and amplifying the DNA) and detecting a polymorphism (comprising obtaining a sample containing human genomic DNA, providing a nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                         ö
restore sensitivity of methicillin resistant Staphylococcus aureus to lactam antibiotic by targeting the molecular mechanism of resistance. The present sequence is used in the exemplification of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human, optineurin, ds, ophthalmological, single nucleotide polymorphism; SNP; glaucoma; progressive ocular hypertensive disorder; glaucoma related disorder; motif; repeat element; regulatory region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New nucleic acid sequences of the optineurin gene are useful to detect polymorphisms particularly single nucleotide polymorphisms in the optineurin promoter to diagnose, prognose and treat glaucoma and related
                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Optineurin promoter motif, repeat element or regulatory region #98
                                                                                                                                                                                                                                                                                                                                                                                         ..
                                                                                                                                                                                                                                                                                                                                                                                    0; Indels
                                                                                                                                                                                                                                                                                        DB 1; Length 10;
                                                                                                                                                                                            Sequence 10 BP; 3 A; 0 C; 7 G; 0 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                         0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 11; SEQ ID NO 100; 159pp; English.
                                                                                                                                                                                                                                                                                        12.3%; Score 9; 1
100.0%; Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                BP.
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AAQ57283/c
ID AAQ57283 standard; mRNA; 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA sequences (AAQ64009-Q64031) are fragments of the 16S rRNA gene from B. bronchiseptica (AAQ55187). The fragments are used as probes to detect porcine atrophic rhinitis caused by the Bordetella bronchiseptica bacterium. Also claimed are 3 DNA fragments complementary to the 436-466 region of the 16S rRNA (AAQ64032-034). A specific DNA seqence from the SI rRNA was selected and 2 probes were designed (AAQ64035 and AAQ64039) for the detection of B. Dronchiseptica. Primers (AAQ64036-37) were used to clone the 16S gene. Sequences (AAQ64034) is the preferred probe used in the detection process. (Updated on 27-AUG-2003 to correct OS field.)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    B.bronchiseptica 16s rRNA fragments - used as probes in the detection porcine atrophic rhinitis.
capable of detecting a SNP located within an optineurin promoter, and detecting the polymorphism). The invention is used to diagnose and prognose glaucoma and also to treat glaucoma related disorders. The present sequence is an optineurin promoter motif, repeat element or putative regulatory region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     detection; procine atrophic rhinitis; hybridisation;
                                                                                                                                                                             Gaps
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                                                                                                                                                                         0; Indels
                                                                                                                                     Query Match
12.3%; Score 9; DB 1; Length 10;
Best Local Similarity 100.0%; Pred. No. 1.3e+03;
Matches 9; Conservative 0; Mismatches 0; Indels
                                                                                                        Seguence 10 BP; 5 A; 1 C; 0 G; 4 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bordetella bronchiseptica; pig raising; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (NISE-) NIHON SEIFUN KK.
(ZENK-) ZENKOKU NOGYO KD RENGOKAI.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 1; Page 10; 12pp; Japanese.
                                                                                                                                                                                                                                                                                                                                 AAQ64023 standard; DNA; 11 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                  (revised)
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bordetella bronchiseptica.
                                                                                                                                                                                                            948 TITAATGTA 956
                                                                                                                                                                                                                                                                                                                                                                                                                                                      16S rRNA gene fragment
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TTTAATGTA 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        rRNA; probe;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               10-JUN-1992;
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22-JUL-1994
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This is a c-myb mRNA target sequence (nucleotide no. 1660) of an enzymatic RNA molecule (ribozyme) which cleaves mRNA associated with the development or maintenance of a restenotic condition. The conon. of the ribozyme necessary to effect a therapeutic treatment is lower than that of an antisense oligonucleotide and the specificity of action is higher. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   prevent
or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                   Specific, cleavage, target RNA, protein, prophylaxis, expression, inhibitor, inhibition, ribozyme, treatment, prevention, psoriasis, asthma, inflammatory diseases, restenosis, cardiovascular condition,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Wound healing; non-MRL healer mouse; quantitative trait locus; QTL; healing response; microsatellite marker; treatment; central nerve; peripheral nerve; nerve injury; SAGE tag; murine; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ô
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Enzymatic RNA molecules which cleave mRNA - used to treat or inflammatory, arthritic, stenotic or cardiovascular diseases
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                                                                                          Enzymatic RNA molecule c-myb mRNA target sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 11 BP; 7 A; 0 C; 3 G; 1 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 3; Page 20; 65pp; English.
                                                                                                                                                                                                                                                                                                 92US-00916763.
92US-00987132.
92US-00989848.
92US-00989849.
93US-00008895.
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                                                                                                                                                               hypertension; arthritis; ss.
                                                                                                                                                                                                                                                                                                                                                                                      (RIBO-) RIBOZYME PHARM INC
                                                                   (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity 100.
Matches 9; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                Draper KG;
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                                                      (revised)
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                                                                                                                                                                                                                     WO9402595-A1,
                                                                                                                                                                                                                                                                                                  17-JUL-1992;
07-DEC-1992;
07-DEC-1992;
07-DEC-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                Sullivan SM,
                                                                                                                                                                                                                                                                         02-JUL-1993;
                                                                                                                                                                                                                                              03-FEB-1994.
                                                    25-MAR-2003
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                                                                 26-JUL-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                conditions.
                                                                                                                                                                                           Synthetic.
                          AAQ57283;
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Gaps

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12.3%; Score 9; DB 1; Length 11; 100.0%; Pred. No. 1.4e+03; ative 0; Mismatches 0; Indels

9; Conservative

Matches

Local Similarity

Query Match

958 CGCTACCAA 966

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The invention provides triplex-forming oligonucleotides (TFO) and their modified derivatives TFO B1-B5 (AAX12862-866) can bind with the promoter region of pre-S gene in inhibition of hepatitis B virus (HBV) adr subtype and TFO B11, B12 and B15 (AAX12868-870) can bind with DR region of HBV.

The oligonucleotides are useful for inhibition of HBV and as drug in treatment of hepatitis B. Since the length of the oligonucleotides can be suitably increased, the stability and specificity of the formed triplex DNA with 2 similar homopoly purine/homopoly pyrimidine fragments are higher. Triplex formed in a Specifically targeting on the HBV gene expression, DNA replication and reproduction, or to produce (DNA) 2:RNA hybrid triplex with target sequence of RNA in stopping RNA reverse transcription, so there is little effect on the human cells. Such oligonucleotides are chemically modified by 3-terminal monophosphorylation, leading to more significant inhibition due to their higher stability, and the degradation products of the modified or 20-MAR-2003 to correct DR field.) (Updated on 27-ANG-2003 to correct OS field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DNA detection; triple helix; identification; bacteria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Triple helix third strand of Hepatitis B virus nucleotides 3151-3161.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                   Triplex-forming oligonuclectides, useful for, e.g. inhibition of hepatitis B virus (HBV).
/*tag= a
/note= "optional monophosphorylation (claim 2)"
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                                                                                                                                                                                                                             (SHAN-) SHANGHAI INST BIOCHEMISTRY CHINESE ACAD.
                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 1, 2; Page 22; 39pp; Chinese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAX14809 standard; DNA; 11 BP.
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                                                                                                                                            98WO-CN000248.
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Best Local Similarity 100.
Matches 9; Conservative
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                                                                                                                                                                                                                                                                                                                    WPI; 1999-288270/27.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                932 CCCTCCTCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
Hepatitis B virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22-DEC-1993;
                                                              WO9920641-A1.
                                                                                                                                                                                         21-OCT-1997;
                                                                                                                                              19-OCT-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This invention describes a novel non-MRL healer mouse (M) having at least one quantitative trait locus selected from those given in the specification, exhibiting an enhanced healing response to a wound compared to mice (m) without the locus. The invention describes a novel method of identifying a gene involved in enhanced wound healing by identifying DNA microsatellite markers which can distinguish healer mice item non-healer mice and identifying microsatellite markers which segregate with enhanced wound healing in progeny of the mice, where a chromosomal locus containing at least one enhanced wound healing gene is identified. A method of treating a wound in a mammal is also disclosed. The new methods are useful for treating wounds, especially central and restoring function after nerve injury in a mammal. (M) is useful as a mammalian model of enhanced wound healing, useful for identifying genes and gene products involved in enhanced wound healing, and to provide methods for wound healing. AAZI861-Z19036 represent murine SAGE Eags from contains mice which are used to illustrate the method of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  .;
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                                                                                                                                                                                                                                                                                                                                                                                  for enhanced wound healing - useful for identifying
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Best Local Similarity 100.0%; Pred. No. 1.4e+03;
Matches 9; Conservative 0; Mismatches 0; Indels
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11
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 13; Page 72; 136pp; English.
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                                                                                                                                                                     98US-0074737P.
98US-0097937P.
98US-0102051P.
                                                                                                                                                                                                                                                                                                                                                                                                             enhanced wound healing genes.
                                                                                                                                 99WO-US002962
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(revised)
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                                                                                                                                                                                                                                                                                                                                                                                       New mammalian model
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Hepatitis B virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Key
misc_feature
                                                                                                                                                                       13-FEB-1998;
26-AUG-1998;
28-SEP-1998;
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20-MAR-2003
28-JUN-1999
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                                                 WO9941364-A2
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        Mus sp.
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Triplex formation; DNA detection; triple helix; identification; bacteria;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Assay of genetic sequences based on triplex formation from double stranded analyte - and hybrid of anchor and reporter sequences, with
                                                                                                                                                                                                                                                                                                                                                                                                                         (PROF-) PROFILE DIAGNOSTIC SCI INC.
                                                                                                                                                                                                                                                                                    AAX14773 standard; DNA; 11 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1999-130384/11.
                                                                                                                                                                                                                                                                                                                                                          Hepatitis B virus
                                                                                                                                                                                                                                                                                                                                                                                               22-DEC-1993;
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The present sequence represents a polynucleotide that is able to form a triple helix with a double stranded sequence. Cytosine bases in the cytosine helix with a double stranded sequence. Cytosine bases in the present can be replaced with 5-methyloytosine for increased triplex stability. The present sequence is used in the assay of the invention, where it can be part of the anchor DNA or reporter DNA sequence. The assay comprises adding a sample containing double-stranded DNA test sequences to an aqueous medium containing at least one complex of anchor DNA, attached to a solid support, and reporter DNA, where either a part of the nanchor DNA or reporter DNA is designed to form a triple-strand structure with part of the test sequence. Triplex formation results in displacement of the reporter DNA which is detected as an indication of the presence of the DNA test sequence. The method is used to detect DNA sequences, particularly for identification of bacteria (by detecting concepnes and Hepatitis B virus
      reporter released if triplex formation occurs, used e.g. to identify
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 11 BP; 0 A; 8 C; 0 G; 3 T; 0 U; 0 Other;
                                                                                                 Disclosure; Col 19-20; 168pp; English.
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Best Local Similarity luv...
9; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present sequence represents a polynucleotide that is able to form a triple helix with a double stranded sequence. Cytosine bases in the present can be replaced with 5-methylcytosine for increased triplex stability. The present sequence is used in the assay of the invention, where it can be part of the anchor DNA or reporter DNA sequence. The assay comprises adding a sample containing double-stranded DNA test sequences to an aqueous medium containing at least one complex of anchor DNA, attached to a solid support, and reporter DNA, where either a part of the anchor DNA or reporter DNA is designed to form a triple-strand structure with part of the test sequence. Triplex formation results in displacement of the reporter DNA which is detected as an indication of the presence of the DNA test sequence. The method is used to detect DNA sequences, particularly for identification of bacteria (by detecting genes for ribosomal RNA) in clinical samples, but also detection of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                         Assay of genetic sequences based on triplex formation from double stranded analyte - and hybrid of anchor and reporter sequences, with reporter released if triplex formation occurs, used e.g. to identify bacteria.
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                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Col 19-20; 168pp; English.
                                                           (PROF-) PROFILE DIAGNOSTIC SCI INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           oncogenes and Hepatitis B virus
92US-00968436,
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                                                                                                                            Hepburn AG, Wang C;
                                                                                                                                                                                           WPI; 1999-130384/11.
29-OCT-1992;
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Gaps .

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Score 9, DB 1; Length 11; Pred. No. 1.4e+03; 0; Mismatches 0; Indels

12.3%; SCOL. 100.0%; Pre

DNA detection; triple helix; identification; bacteria; Triple helix third strand of Hepatitis B virus nucleotides 1614-1624. Assay of genetic sequences based on triplex formation from double stranded analyte - and hybrid of anchor and reporter sequences, with reporter released if triplex formation occurs, used e.g. to identify Disclosure; Col 17-18; 168pp; English. (PROF-) PROFILE DIAGNOSTIC SCI INC. AAX14747 standard; DNA; 11 BP. 93US-00173489. 92US-00968436. 24-MAR-1999 (first entry) oncogene; virus; ss. Hepburn AG, Wang C; WPI; 1999-130384/11. Triplex formation; Homo sapiens 22-DEC-1993; 29-OCT-1992; JS5861244-A. 19-JAN-1999. Synthetic.

The present sequence represents a polynucleotide that is able to form a triple helix with a double stranded sequence. Cytosine bases in the present can be replaced with 5-methyloytosine for increased triplex stability. The present sequence is used in the assay of the invention,

93US-00173489. 92US-00968436, ô

Gaps

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Indels

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Mismatches

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9; Conservative

Matches

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The invention relates to identifying (M1) genes in vitro that, in humans or animals, are important for skin ageing and/or skin stress by serial analysis of gene expression between mixtures of transcribed and optionally translated, genetically encoded factors (A) obtained from young and aged skin, to identify that genes that show strong differential useful for: identifying maxkers of skin ageing and/or stress; determining skin ageing and/or stress; and identifying or determining the effects of sharmaceutical and or comentic agents for control of skin ageing. The present sequence is one of a group of human skin ageing/stress related expressed sequence tags (ABQ86246-ABQ87680) of the invention
where it can be part of the anchor DNA or reporter DNA sequence. The assay comprises adding a sample containing double-stranded DNA test sequences to an aqueous medium containing at least one complex of anchor DNA, attached to a solid support, and reporter DNA, where either a part of the anchor DNA or reporter DNA is designed to form a triple-strand structure with part of the test sequence. Triplex formation results in displacement of the reporter DNA which is detected as an indication of the presence of the DNA test sequence. The method is used to detect DNA sequences, particularly for identification of bacteria (by detecting genes for ribosomal RNA) in clinical samples, but also detection of
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                                                                                                                                                                                                                                                                     12.3%; Score 9; DB 1; Length 11; 100.0%; Pred. No. 1.4e+03; tive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                 Sequence 11 BP; 0 A; 8 C; 0 G; 3 T; 0 U; 0 Other;
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                                                                                                                                                                                               oncogenes and Hepatitis B virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ABQ86582 standard; cDNA; 11 BP.
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Best Local Similarity
Matches 9; Conserv
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ABQ86582/
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12.3%; Score 9; DB 1; Length 11; 100.0%; Pred. No. 1.4e+03;

Query Match Best Local Similarity

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The invention relates to identifying (M1) genes in vitro that, in humans or animals, are important for skin ageing and/or skin stress by serial analysis of gene expression between mixtures of transcribed and optionally translated, genetically encoded factors (A) obtained from young and aged skin, to identify that genes that show strong differential useful for: identifying markers of skin ageing and/or stress; determining skin ageing and/or stress; and identifying or determining the effects of pharmacoustical agents for control of skin ageing. The present sequence is one of a group of human skin ageing/stress related expressed sequence tags (ABQ86246-ABQ87680) of the invention
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Identifying genes involved in skin stress and aging, useful e.g. in screening for cosmetic or therapeutic agents, based on differential gene
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                                                                                                                                                                                                                    tag;
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                                                                                                                                                                                    Human skin stress/ageing related EST SEQ ID NO 962.
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ABV65978 standard; cDNA; 11
                                                                                       7207/c
ABQ87207 standard; cDNA; 11
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908 TITICITIG 916
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                                                                      RESULT 2574
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ID ABV65
XX
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The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically seconded from Skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression.

(M1) is useful for identifying genes involved in skin homeostasis; to promotes skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitis; sunburn; psoriasis; scleroderma; ichthyosis; atopic dermatitis; acne; seborrhea; lupus erythematosus; rosacea; melanoma; basal cell carcinoma; and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag (EST) of the invention
                                 Human, skin, dermatological, vulnerary, antipsoriatic, antiseborrhaeic, immunosuppressive, antiinflammatory, cytostatic, SAGE, neurodermatitis, psoriasis, dermatitis, skin cancer, EST; expressed sequence tag, ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             In vitro identification of skin-expressed genes, useful for determining homeostasis and identifying cosmetic or pharmaceutical agents against
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immunosuppressive, antiinflammatory, cytostatic, SAGE, neurodermatitis,
psoriasis, dermatitis, skin cancer, EST, expressed sequence tag, ss.
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ABV71682 standard; cDNA; 11 BP.
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ABV71682/C
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Hofmann K;

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The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression.

(M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitis; sunburn; psoriasis; scleroderma; ichthyosis; atopic dermaitis; acne; seborrhea; lupus erythematosus; rosacea; melanoma; basal cell carcinoma; and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag (ESI) of the invention
                                                                                                                            In vitro identification of skin-expressed genes, useful for determining homeostasis and identifying cosmetic or pharmaceutical agents against
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       In vitro identification of skin-expressed genes, useful for determining homeostagis and identifying cosmetic or pharmaceutical agents against e.g. skin cancer.
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                                                              Hofmann K;
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03-JAN-2001; 2001DE-01000127
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                                                              Conradt M,
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                                                              Petersohn D,
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DB 1; Length 11;

12.3%; Score 9; DB 1; Ler 100.0%; Pred. No. 1.4e+03; tive 0; Mismatches 0;

920

Disclosure; Page 72; 1345pp; German

20-DEC-2001; 2001WO-EP015179.

WO200253774-A2

11-JUL-2002

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 Length 11;
 DB 1; Len
Query Match 12.3%; Score 9; DB 1
Best Local Similarity 100.0%; Pred. No. 1.4
Matches 9; Conservative 0; Mismatches
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945 TGGTTTAAT 953 ò 셤

The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (\$AGE) so as to identify skin-expressed genes and quantify their expression (\$AGE) (M1) is useful for identifying genes involved in skin homeostasis, to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitis; sumburn; psoriasis; scleroderma; ichthyosis, atopic dermatitis; acnes; seborrhea; lupus ertytematosus; rosacea; melanoma; basal cell carcinoma, and carcinoma or sarcoma of the form of the present sequence is that of a human expressed sequence tag

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ABV70743;

21-OCT-2002 (first entry)

Human skin EST 8529

WO200253774-A2

20-DEC-2001; 2001WO-EP015179.

03-JAN-2001; 2001DE-01000127

WPI; 2002-590638/63

Sequence 11 BP; 5 A; 2 C; 1 G; 3 T; 0 U; 0 Other;

Gaps ; 0 12.3%; Score 9; DB 1; Length 11; 100.0%; Pred. No. 1.4e+03; tive 0; Mismatches 0; Indels

947 GITTAATGT 955

The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from Skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression. (M1) is useful for identifying genes involved in skin homeostasis; to promotes skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitis; sunburn; psoriasis; scleroderma; ichthyosis; atopic dermatitis; acne; seborrhea; lupus erythematosus; rosacea, melanoma; basal cell carcinoma; and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag

In vitro identification of skin-expressed genes, useful for determining homeostasis and identifying cosmetic or pharmaceutical agents against e.g. skin cancer.

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Hofmann

Petersohn D, Conradt M,

(HENK) HENKEL KGAA

WPI; 2002-590638/63.

20-DEC-2001; 2001WO-EP015179. 03-JAN-2001; 2001DE-01000127

WO200253774-A2. Homo sapiens

11-JUL-2002

Disclosure; Page 118; 1345pp; German.

Sequence 11 BP; 5 A; 2 C; 2 G; 2 T; 0 U; 0 Other;

X S

10 TGGTTTAAT 2

RESULT 2579

ABV70743/c ID ABV70743 standard; cDNA; 11

Gaps ö

12.3%; Score 9; DB 1; Length 11; 100.0%; Pred. No. 1.4e+03; trive 0; Mismatches 0; Indels

Local Similarity 100. nes 9; Conservative

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Query Match

912 CTTTGGTCT 920 CTTTGGTCT 11

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Sequence 11 BP; 0 A; 2 C; 4 G; 5 T; 0 U; 0 Other;

Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.

Homo sapiens

ABV65564 standard; cDNA; 11 BP.

RESULT 2578

ABV65564/c

(HENK) HENKEL KGAA

Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.

Human skin EST 3350

21-OCT-2002

ABV65564;

Petersohn D, Conradt M, Hofmann K;

In vitro identification of skin-expressed genes, useful for determining homeostasis and identifying cosmetic or pharmaceutical agents against e.g. skin cancer.

Claim 24; Page 273; 1345pp; German.

The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression. (M1) is useful for identifying genes involved in skin homeostasis; to promotes skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitis; sunburn; psoriasis; scleroderma; ichthyosis; atopic dermatitis; acne; seborrhea; lupus erythematosus; rosacea, melanoma; basal cell carcinoma; and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag (EST) of the invention

Query Match
Best Local Similarity 100...
9; Conservative

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11

RESULT 2580

Hofmann K;

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In vitro identification of skin-expressed genes, useful for determining homeostasis and identifying cosmetic or pharmaceutical agents against e.g. skin cancer.
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                                                                         20-DEC-2001; 2001WO-EP015179.
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     Homo sapiens.
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                                                                                                          Human, skin, dermatological, vulnerary, antipsoriatic, antiseborrhaeic, immunosuppressive, antiinflammatory, cytostatic, SAGB, neurodermatitis, psoriasis, dermatitis, skin cancer, EST, expressed sequence tag, ss.
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immunosuppressive, antiinflammatory, cytostatic, SAGE, neurodermatitis,
psoriasis, dermatitis, skin cancer, EST, expressed sequence tag, ss.
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             ABV71325 standard; cDNA; 11 BP
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                                    ABV71325;
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The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression.

(M1) is useful for identifying genes involved in skin homeostasis; to promotes skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitis; sunburn, psoriaeis, scleroderma; ichthyosis; atopic dermatitis; acne; seborrhea; lupus erythematosus; rosacea; melanoma; basal cell carcinoma; and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag
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The invention relates to in vitro identification (MI) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from Skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression.

(MI) is useful for identifying genes involved in skin homeostasis, to promotes skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitis; sunburn, psoriasis, scleroderma; ichthyosis; atopic dermatitis; acne; seborrhea; lupus erythematosus; ichthyosis; metanoma; basal cell carcinoma, and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression. In vitro identification of skin-expressed genes, useful for determining homeostasis and identifying cosmetic or pharmaceutical agents against e.g. skin cancer. Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss. In vitro identification of skin-expressed genes, useful for determining homeostasis and identifying cosmetic or pharmaceutical agents against Gaps ö 0; Indels DB 1; Length 11; . 1.4e+03; Sequence 11 BP; 8 A; 1 C; 2 G; 0 T; 0 U; 0 Other; 12.3%; Scort 100.0%; Pred. No. 1... 0; Mismatches Disclosure; Page 167; 1345pp; German. Hofmann K; Claim 24; Page 294; 1345pp; German. 20-DEC-2001; 2001WO-EP015179. 03-JAN-2001; 2001DE-01000127. ABV71372 standard; cDNA; 11 (first entry) Petersohn D, Conradt M, Similarity 100. 9; Conservative 908 TITICITIG 916 10 TTTTCTTTG 2 Human skin EST 9158. (HENK) HENKEL KGAA. WPI; 2002-590638/63 WPI; 2002-590638/63 e.g. skin cancer. WO200253774-A2 Homo sapiens 21-OCT-2002 11-JUL-2002. Query Match Best Local S ABV71372; RESULT 2583 ઠે 요

ö The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression. (M1) is useful for identifying genes involved in skin homeostasis, to promotes skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitis; sunburn; psoriaals; scleroderma; ichthyosis; atopic dermatitis; acne; seborrhea; lupus erythematosus; icothnyosis; atopic dermatitis; acne; seborrhea; lupus erythematosus; scin. The present sequence is that of a human expressed sequence tag (EST) of the invention In vitro identification of skin-expressed genes, useful for determining homeostasis and identifying cosmetic or pharmaceutical agents against Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGB; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss. Gaps ; 0 12.3%; Score 9; DB 1; Length 11; 100.0%; Pred. No. 1.4e+03; Sequence 11 BP; 4 A; 2 C; 5 G; 0 T; 0 U; 0 Other; Score 9; DB 1; Ler ; Pred. No. 1.4e+03; 0; Mismatches 0; ĸ Claim 24; Page 280; 1345pp; German. Hofmann 20-DEC-2001; 2001WO-EP015179. 03-JAN-2001; 2001DE-01000127. ABV70941 standard; cDNA; 11 (first entry) Query Match
Best Local Similarity 100.
Matches 9; Conservative Σ Conradt 3 CTTTGGTCT 11 WPI; 2002-590638/63. (HENK) HENKEL KGAA Human skin EST 8727 Query Match Best Local Similarity 912 CITIGGICT e.g. skin cancer. WO200253774-A2. 'n 21-OCT-2002 11-JUL-2002. Petersohn ABV70941; RESULT 2584 ABV70941/c Homo 셤 à

(M1) is useful for identifying genes involved in skin homeostasis; to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitis; sunburn; psoriaais; scleroderma; ichthyosis; atopic dermatitis; acne; seborrhea; lupus erythematosus; rosacea; melanoma; basal cell carcinoma; and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag (EST) of the invention

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Length 11;

Sequence 11 BP; 0 A; 2 C; 4 G; 5 T; 0 U; 0 Other;

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The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression. (M1) is useful for identifying genes involved in skin homeostasis to determine skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitis; sunburn; psoriasis, scleroderma; ichthyosis; atopic dermatitis; acne; seborrhea; lupus erythematosus; rosacea, melanoma; basal cell carcinoma, and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag
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21-OCT-2002 (first entry)
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                                        Human skin EST 1306
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                                                                                 cerecter 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 2002-590638/63.
                                                                                                                                                                                                                                                                                                                    Human skin EST 1108
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 Homo sapiens.
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                                                                                                                                                                                                                                 ABV63322;
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ABV63520/c
ID ABV6352(
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AC ABV6352(
XX
  Matches
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Matches
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e.g. skin cancer.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to scrial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression. (M1) is useful for identifying genes involved in skin homeostasis or promotes skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitis; sunburn, psoriasis, scleroderma; icthhyosis; atopic dermatitis; acne; seborrhea; lupus erythematosus; rosacea; melanoma; basal cell carcinoma; and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag (BST) of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human; skin; dermatological; vulnerary; antipsoriatic; antiseborrhaeic; immunosuppressive; antiinflammatory; cytostatic; SAGE; neurodermatitis; psoriasis; dermatitis; skin cancer; EST; expressed sequence tag; ss.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        In vitro identification of skin-expressed genes, useful for determining homeostasis and identifying cosmetic or pharmaceutical agents against
                                                                                                                                                                                                                                                                                                          In vitro identification of skin-expressed genes, useful for determining homeostasis and identifying cosmetic or pharmaceutical agents against
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 11 BP; 8 A; 1 C; 2 G; 0 T; 0 U; 0 Other;
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                                                                                                                                                                                                   Hofmann K;
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                                20-DEC-2001; 2001WO-EP015179
                                                                                        03-JAN-2001; 2001DE-01000127
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Best Local Similarity 100.
Matches 9; Conservative
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                                                                                                                                                                                                   Conradt M,
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                                                                                                                                           (HENK ) HENKEL KGAA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             21-OCT-2002
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                              The invention relates to in vitro identification (M1) of genes expressed in the skin of humans or animals by subjecting a mixture of genetically encoded factors from skin, to serial analysis of gene expression (SAGE) so as to identify skin-expressed genes and quantify their expression. (M1) is useful for identifying genes involved in skin homeostasis; to promotes skin homeostasis and to test agent (A) that maintains or promotes skin homeostasis or that can be used for treating skin disorders, specifically neurodermatitis; sunburn; psoriaais, scleroderma; ichthyosis; atopic dermatitis; acne; seborrhea; lupus erythematosus; rosacea, melanoma; basal cell carcinoma; and carcinoma or sarcoma of the skin. The present sequence is that of a human expressed sequence tag (EST) of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The present sequence represents a fragment of the porcine circovirus genome associated with MAP. MAP is the french acronym for piglet fatal wasting disease. The polypetides can be used to detect anti-MAP antibodies. The antibodies can be used to detect MAP antigens. The nucleotide sequences can be used as probes or primers for detecting MAP nucleic acids. The nucleotide sequences, polypeptides, vectors, (pseudo)viral particles, transformed cells and compounds selected by the screening assay can be used in pharmaceutical compositions. The polypeptides, nucleotide sequences, vectors and transformed cells can be used in vaccines against MAP circovirus infection. The vectors, (pseudo)viral particles and transformed cells can be used for gene
                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            porcine circovirus MAP - useful in vaccines infection and in gene therapy.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MAP; piglet fatal wasting disease; vaccine; circovirus infection;
gene therapy; ss.
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                                                                                                                                                                                                                                                                                                                                                                              Length 11;
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                                                                                                                                                                                                                                                                                                                                   Sequence 11 BP; 4 A; 0 C; 6 G; 1 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                                                                                         Query Match
12.3%; Score 9; DB 1; Lei
Best Local Similarity 100.0%; Pred. No. 1.4e+03;
Matches 9; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Fragment of the porcine circovirus genome.
Disclosure; Page 200; 1345pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Le Cann P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 5; Page 59; 89pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAX85598 standard; DNA; 12 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Nucleotide sequence of against MAP circovirus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  935 TCCTCTTCA 943
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            11 recrerrea 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Porcine circovirus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           05-DEC-1997;
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Best Local Similarity

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Sequence 12 BP; 0 A; 5 C; 1 G; 6 T; 0 U; 0 Other;
                                                                                                                                                                                                                                                                                                                   Sequence 12 BP; 1 A; 4 C; 3 G; 4 T; 0 U; 0 Other;
                                                                                                                                                                               Kalbfleisch TS,
                                                                                    (first entry)
                          Conservative
                                   CCTCCTCTT 941
                                           CCTCCTCTT 12
                                                                                                                                                                       (CURA-) CURAGEN CORP.
                                                                                                                                                                                        WPI; 2000-349567/30.
                     Local Similarity
                                                                                                                                                                                                                                                                                                           sequences used
                                                                                   04-SEP-2000
                                                                                                                                                     13-JUN-1997;
                                                                                                                                                              14-JUN-1996;
                                                                                                                                    US6057101-A.
                                                                                                                                            02-MAY-2000.
                                                                                                                                                                                                               interaction.
                                                                                                                                                                               Knight JR,
                                                                                                                           Synthetic.
                                   933
                                                                          AAA55929;
                 Query Match
                                                         2590
                          Matches
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The present invention describes a method for detecting (D) at least 1 protein-protein interaction (PPI) by recombinantly expressing within a protein-protein interaction (PPI) by recombinantly expressing within a population of host cells, populations of first and second fusion proteins comprising DNA binding domain (DBD) and transcriptional regulatory domain (TRD) respectively and detecting the regulation of transcription of mucleotide sequence of host cells operably linked to a promoter driven by DBD. The detection method (D) is useful for identifying inhibitors of PPI for therapeutic use, and for detecting specific cell types, tissue types, cragge of development and disease states. From the population of the proteins characteristic of the particular tissue or a cell-type, all possible detectable PPI that occur can be identified and genes encoding these proteins can be isolated. Thus, parallel analysis of two cell types enumerates PPI that are common to both and those that are specific to both. This analysis has significant value since PPI specific to a disease can also be isolated in rapid fashion. The number of false positives and can also be isolated in rapid fashion. The number of false positives and common the proteins of the particular and proteins can also be isolated in rapid fashion. The number of false positives and common the proteins of the particular content of the particular content content content of the particular content 
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                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Yeast; detection; protein-protein interaction; DNA-binding domain; characterisation; dentification; protein pathway information; protein interaction domain; screening; PCR primer; adapter; linker; fusion protein; inhibitor; regulation; ss.
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12.3%; Score 9; DB 1; Length 12; 100.0%; Pred. No. 1.4e+03; trive 0; Mismatches 0; Indels
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12.3%; Score 9; DB 1; Length 12;

Query Match

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      genes encoding proteins that inceract with each other, via the reconstitution of a transcription factor and hence reporter gene activation. Proteins are fused to either the yeast DNA-binding domain of a transcriptional activator or to the activation domain of a transcriptional activator. The present sequence is a linker used in the present invention as an adapter in the analysis of yeast fusion genes. The present method may be used to identify protein-protein interactions and genes encoding the interacting proteins relevant to a particular tissue, stage or disease e.g. cancer
                                                                                                                                                                                                                                                                     Linker; yeast; two-hybrid system; protein-protein interaction; cancer;
                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
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                           Indels
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            Pred. No. 1.4e+03;
; Mismatches 0;
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100.0%; Pre
                                                                                                                                                       AAA73441 standard; DNA; 12 BP.
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                                                                                                                                                                                                               (first entry)
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                         9; Conservative
                                                                                                                                                                                                                                                                                                               cerevisiae.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                                                                                                                                                                                                                           Claim 1; SEQ ID NO 290772; 29pp + Sequence Listing; German.
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Best Local Similarity 100.0%; Pred. No. 1.4e+03;
Matches 9; Conservative 0; Mismatches 0; Indels
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                                                           SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                           Oligonucleotide primer SEQ ID NO 295709 for detecting SNP TSC0016695.
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Matches 9; Conserv
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ABH90779

RESULT 2593

Query Match

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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nuclectide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but five.wipo.int/pub/published_pct_sequences
                          Claim 1; SEQ ID NO 350587; 29pp + Sequence Listing; German.
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ABI50922 standard; DNA; 12 BP. RESULT 2595

Oligonucleotide primer SEQ ID NO 350895 for detecting SNP TSC0046961. 22-FEB-2002 (first entry) ABI50922;

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Homo sapiens

WO200177384-A2

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06-APR-2001; 2001WO-IB000713.

07-APR-2000; 2000DE-01019173

(EPIG-) EPIGENOMICS AG

Berlin K; Olek A, Piepenbrock C,

WPI; 2001-657177/75.

Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine methylation status.

Claim 1; SEQ ID NO 350895; 29pp + Sequence Listing; German.

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABF59989, ABF00010-ABF99989, ABH00010-ABF99989, ABH00010-ABF99989, ABH00010-ABF99989, ABH00010-ABF99980 and ABI00010-ABF99980 data for this patent did not form part of the printed specification, but

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABT00010-ABI82073 trepresent the oligomers described in the invention. NoTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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central nervous system; gastrointestinal; respiratory; immune; metabolic
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                                                                                                                                                                                           SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                               Oligonuclectide primer SEQ ID NO 374505 for detecting SNP TSC0060748.
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                                                        ABI74532 standard; DNA; 12
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Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine methylation status.
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              WPI; 2001-657177/75.
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                              designed to detect methylation status.
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and cytosine methylation status in chemically pretreated genomic DNA. The oligomucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, contral nervous system, cardiovascular and metabolic disorders. The oligomers a slso used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF99899 and ABI00010-ABF32073 represent the oligomers described in the invention. NOTE: The sequence was obtained in electronic format from WIPO at the printed specification, but the wipo.int/pub/published_pct_sequences
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12.3%; Score 9; DB 1; Les
Best Local Similarity 100.0%; Pred. No. 1.4e+03;
Matches 9; Conservative 0; Mismatches 0;
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Seguence 12 BP; 5 A; 0 C; 2 G; 5 T; 0 U; 0 Other;

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP)

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99889, ABR00010-ABF99889, ABH00010-ABH99889 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic form at from NIPO at
                                                                                               SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                 Oligonucleotide primer SEQ ID NO 274705 for detecting SNP TSC0003650.
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                                                                                                                                                                                                                                                                                                                                  SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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12.3%; Score 9; DB 1; Length 12; 100.0%; Pred. No. 1.4e+03; tive 0; Mismatches 0; Indels
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                                                         Query Match
12.3%; Score 9; DB 1; Length 12;
Best Local Similarity 100.0%; Pred. No. 1.4e+03;
Matches 9; Conservative 0; Mismatches 0; Indels
                             Sequence 12 BP; 3 A; 1 C; 0 G; 8 T; 0 U; 0 Other;
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ABH74720 standard; DNA; 12

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Berlin K;

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting call type differentiation. ABC00010-ABC99989, ABF00010-ABE99989 and ABI00010-ABI82073. represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                         Claim 1; SEQ ID NO 326719; 29pp + Sequence Listing; German.
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Berlin K;

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but ftp.wipo.int/pub/published_pct_sequences
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Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                Claim 1; SEQ ID NO 301657; 29pp + Sequence Listing; German.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SNP, single nuclectide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the Oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Oligonucleotide primer SEQ ID NO 304566 for detecting SNP TSC0020999.
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                                                                                                                                           12.3%; Score 9; DB 1; Length 12; 100.0%; Pred. No. 1.4e+03; tive 0; Mismatches 0; Indels
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                                                                                                           Sequence 12 BP; 1 A; 2 C; 0 G; 9 T; 0 U; 0 Other;
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic discorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABE09989, ABF00010-ABE99989, ABF00010-ABE99989 and ABI00010-ABE32073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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WPI; 2001-657177/75.
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Query Match 12.3%; Score 9; DB 1; Length 12; Best Local Similarity 100.0%; Pred. No. 1.4e+03; Matches 9; Conservative 0; Mismatches 0; Indels

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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                               Claim 1; SEQ ID NO 286420; 29pp + Sequence Listing; German.
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                                                                                                                                                         (EPIG-) EPIGENOMICS AG
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                                                     Homo sapiens
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genemic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genemic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, contral nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at

Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine

Berlin

Piepenbrock C,

olek A,

WPI; 2001-657177/75.

methylation status.

07-APR-2000; 2000DE-01019173.

(EPIG-) EPIGENOMICS AG

Claim 1; SEQ ID NO 345878; 29pp + Sequence Listing; German.

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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                Oligonucleotide primer SEQ ID NO 345878 for detecting SNP TSC0044262
                                                                             ABI45905 standard; DNA; 12 BP.
                                                                                                                         22-FEB-2002 (first entry)
934 CICCICITC 942
         AB145905,
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                  Oligonucleotide primer SEQ ID NO 348250 for detecting SNP TSC0045503.
                                                                                                                                                                                                                                                                                                                                                                                                   ligonucleotides, useful for diagnosis and cell typing, is to detect single-nucleotide polymorphisms and cytosine
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                                                            ABI48277 standard; DNA; 12 BP.
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                                    RESULT 2611
ABI48277
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Claim 1; SEQ ID NO 348250; 29pp + Sequence Listing; German.

06-APR-2001; 2001WO-IB000713.

WO200177384-A2.

18-OCT-2001

Homo sapiens

Sequence 12 BP; 6 A; 0 C; 3 G; 3 T; 0 U; 0 Other;

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99889, ABC0010-ABE99889, ABC0010-ABE99899 and ABI00010-ABE82073 teprement the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but two obtained in electronic format from WIPO at
            This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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12.3%; Score 9; DB 1; Length 12; 100.0%; Pred. No. 1.4e+03; tive 0; Mismatches 0; Indels
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RESULT 2614

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Olek A, Piepenbrock C, Berlin K;
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                                                                                                        SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                   Oligonucleotide primer SEQ ID NO 357261 for detecting SNP TSC0050537.
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               ABI57288 standard; DNA; 12 BP.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                                                                                                                                                                                                                                                                                                 Berlin K;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                           Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                      Claim 1; SEQ ID NO 377041; 29pp + Sequence Listing; German.
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             WPI; 2001-657177/75
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range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Similarity 100.0%; Pred. No. 1.4e+03;
9; Conservative 0; Mismatches 0;
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12.3%; Score 9; DB 1; Length 12; 100.0%; Pred. No. 1.4e+03;

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at
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                                                                             SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                      Oligonucleotide primer SEQ ID NO 278551 for detecting SNP TSC0006119
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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22-FEB-2002
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                                                                                                                                                                                                                                                                                                                                                                SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
  Gaps
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ss 0; Indels
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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ABI04695 standard; DNA; 12
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                                                                    This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99889, ABF00010-ABF99899 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but two but into int/pub/published_pot_sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                     Claim 1; SEQ ID NO 304668; 29pp + Sequence Listing; German.
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Local Similarity 100.0%; Pred. No. 1.4e+03;
nes 9; Conservative 0; Mismatches 0;
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methylation status.
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                             ABI15017 standard; DNA; 12 BP.
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Matches 9; Conservative
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data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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(EPIG-) EPIGENOMICS AG.

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nucleic acid; cytosine methylation; cardiovascular; primer; ss; nervous system; gastrointestinal; respiratory; immune; metabolic.
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                               Olek A, Piepenbrock C,
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acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and range of diseases including immune system, gastrointestinal, respiratory. Central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC09989, ABC0010-ABC99899 ABN0010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form par of the printed specification, but was obtained in electronic format from WIPO at this printed but is but int/pub/published_pct_sequences
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ABIS 9942/C

ABIS 9
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Seguence 12 BP; 5 A; 2 C; 0 G; 5 T; 0 U; 0 Other;

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                                                                                                                                                                                                                                                                                                                                                       SNP; single nucleotide polymorphism; human; diagnosis; FNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                        Oligonucleotide primer SEQ ID NO 381690 for detecting SNP TSC0064487.
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  DB 1; Length 12;
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Query Match
12.3%; Score 9; DB 1; Ler
Best Local Similarity 100.0%; Pred. No. 1.4e+03;
Matches 9; Conservative 0; Mismatches 0;
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ID ABI19671 standard; DNA; 12 BP.
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Matches 9; Conservative
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                                                                            948 TTTAATGTA 956
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 therefresh the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                                                                                                                                                                                                                Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; SEQ ID NO 304151; 29pp + Sequence Listing; German.
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                                                                                                                                                                                   (EPIG-) EPIGENOMICS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nuclectide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99889, ABC0010-ABH99889 and ABI00110-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form par of the printed specification, but the was obtained in electronic format from WIPO at.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                        SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                Oligonucleotide primer SEQ ID NO 319644 for detecting SNP TSC0029341.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        designed to detect methylation status.
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ABI39603 standard; DNA; 12 BP.
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Set of oligonuclectides, useful for diagnosis and cell typing, : designed to detect single-nucleotide polymorphisms and cytosine
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                                                                                                                                                Claim 1; SEQ ID NO 304565; 29pp + Sequence Listing; German.
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                                                                            methylation status.
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Best Local Similarity
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
Oligonucleotide primer SEQ ID NO 339576 for detecting SNP TSC0041078.
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Homo sapiens

WO200177384-A2.

07-APR-2000; 2000DE-01019173

(EPIG-) EPIGENOMICS AG

Berlin Piepenbrock C, WPI; 2001-657177/75. Claim 1; SEQ ID NO 339576; 29pp + Sequence Listing; German.

methylation status.

Set of oligonucleotides, useful for diagnosis and cell typing, : designed to detect single-nucleotide polymorphisms and cytosine

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The

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Gaps

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12.3%; Score 9; DB 1; Length 12; 100.0%; Pred. No. 1.4e+03; ative 0; Mismatches 0; Indels

Best Local Similarity 100. Matches 9; Conservative

Query Match

Sequence 12 BP; 5 A; 0 C; 3 G; 4 T; 0 U; 0 Other;

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oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at from thy pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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designed to detect single-nucleotide polymorphisms and cytosine
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les 9; Conservative 0; Mismatches
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Oligonucleotide primer SEQ ID NO 353877 for detecting SNP TSC0048780.
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                                                                                                                                                                                                                                                                                                                              SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                     Oligonucleotide primer SEQ ID NO 291024 for detecting SNP TSC0014601.
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designed to detect single-nucleotide polymorphisms and cytosine
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABE9989, ABF00010-ABE9989, ABF00010-ABE9989, and ABI00010-ABE82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but twipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                              oligonucleotides, useful for diagnosis and cell typing, is to detect single-nucleotide polymorphisms and cytosine
                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 1; SEQ ID NO 364347; 29pp + Sequence Listing; German.
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methylation status.
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Query Match
12.3%; Score 9; DB 1; Length 12;
Best Local Similarity 100.0%; Pred. No. 1.4e+03;
Matches 9; Conservative 0; Mismatches 0; Indels
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Oligonucleotide primer SEQ ID NO 267920 for detecting SNP TSC0000690.

SNP; single nucleotide polymorphism; human; diagnosis; FNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Berlin K;

WPI; 2001-657177/75

Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.

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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABE99989, ABF00010-ABE99989, ABF00010-ABE99989, and ABI00010-ABI32073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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Claim 1; SEQ ID NO 267920; 29pp + Sequence Listing; German.
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic. Oligonucleotide primer SEQ ID NO 271891 for detecting SNP TSC0002645. 22-FEB-2002 (first entry)

ABH71914 standard; DNA; 12 BP.

ABH71914;

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Homo sapiens.

WO200177384-A2.

18-OCT-2001.

06-APR-2001; 2001WO-IB000713.

07-APR-2000; 2000DE-01019173

(EPIG-) EPIGENOMICS AG

Berlin K; Olek A, Piepenbrock C,

WPI; 2001-657177/75.

Set of oligonucleotides, useful for diagnosis and cell designed to detect single-nucleotide polymorphisms and methylation status.

Claim 1; SEQ ID NO 271891; 29pp + Sequence Listing; German.

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF9989, ABH00010-ABH9989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at

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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                            Oligonucleotide primer SEQ ID NO 279863 for detecting SNP TSC0007882.
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                            ABH79870 standard; DNA; 12
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RESULT 2642
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                                                                                                                                                                                                                                                                                                                           SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Best Local Similarity 100.
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Matches 9; Conservative
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Berlin K;

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from "MPD at ftp.wipo.int/pub/published_pct_sequences
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, contral nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 tepresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                          range of diseases including immune system, gastrointestinal, respiratory, cardiovascular and metabolic disorders. The colisomers are also used for detecting cell type differentiation. ABC0010-ABC0989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the olisomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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   oligonucleotides are used for diagnosis and/or prognosis of cancer and a
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                                                                                                                                                                                                                                                                                    SND; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                         Oligonucleotide primer SEQ ID NO 370762 for detecting SNP TSC0058378.
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ive 0; Mismatches 0; Indels
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                                                     943 ATTGGTTTA
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                                                              SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                       Oligonucleotide primer SEQ ID NO 360790 for detecting SNP TSC0052292.
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo int/pub/published_pct_sequences
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                                                                                                                                set or oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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Berlin K;
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                                                                                                               This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010 ABC99989, ABF00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at the printed specification, but fire wipo.int/pub/published_pct_sequences
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designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                        Claim 1; SEQ ID NO 376227; 29pp + Sequence Listing; German.
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represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PMA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                              SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                              Oligonucleotide primer SEQ ID NO 270293 for detecting SNP TSC0002077.
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11D ABH70316

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MAC ABH7
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 targressent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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designed to detect single-nucleotide polymorphisms and cytosine
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                                                                                                                                        This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABE99989, ABF00010-ABE99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but fur wipo.int/pub/published_pct_sequences
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designed to detect single-nucleotide polymorphisms and cytosine
                                                                                                                     Claim 1; SEQ ID NO 278586; 29pp + Sequence Listing; German.
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12.3%; Score 9; DB 1; Length 12;
Best Local Similarity 100.0%; Pred. No. 1.4e+03;
Matches 9; Conservative 0; Mismatches 0; Indels
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Claim 1; SEQ ID NO 304747; 29pp + Sequence Listing; German.

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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12.3%; Score 9; DB 1; Length 12;
Best Local Similarity 100.0%; Pred. No. 1.4e+03;
Matches 9; Conservative 0; Mismatches 0; Indels
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, contral nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at
                                                                                                                                                                 SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                            Oligonucleotide primer SEQ ID NO 290115 for detecting SNP TSC0014219.
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designed to detect single-nucleotide polymorphisms and cytosine
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ABH90122 standard; DNA; 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             SNP, single nuclectide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                       12.3%; Score 9; DB 1; Length 12; 100.0%; Pred. No. 1.4e+03; tive 0; Mismatches 0; Indels
BP; 3 A; 0 C; 3 G; 6 T; 0 U; 0 Other;
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from MIPO at
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                   Claim 1; SEQ ID NO 357902; 29pp + Sequence Listing; German.
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Matches 9; Conservative 0; Mismatches 0; Indels
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central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010 -ABC9989, ABF00010-ABF9989, ABF00010-ABF9989, ABF00010-ABF9989, ABF00010-ABF9989 and ABI00100-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this parent did not form part of the printed specification, but was obtained in electronic format from WIPO at thick pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                       SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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12.3%; Score 9; DB 1; Length 12;
Best Local Similarity 100.0%; Pred. No. 1.4e+03;
Matches 9; Conservative 0; Mismatches 0; Indels
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Oligonucleotide primer SEQ ID NO 274779 for detecting SNP TSC0003673.
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Matches 9; Conservative
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948 TITAATGTA 956
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06-APR-2001; 2001WO-IB000713.

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, contral nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 trepresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                     SND; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                 Oligonucleotide primer SEQ ID NO 278939 for detecting SNP TSC0006650.
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                                      SNP; single nucleotide peptide nucleic acta
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RESULT 2666 ABI08267

Matches

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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99999, ABF00010-ABF99999, ABH00010-ABH99999 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                        This invention describes novel oligonucleotide primers or peptide nucleic acid (PMA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99899, ABH00010-ABH99989 and ABI00010-ABH82073 tepresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but fep.wipo.int/pub/published_pct_sequences
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Claim 1; SEQ ID NO 340600; 29pp + Sequence Listing; German.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                            DB 1; Length 12; 1.4e+03;
                                                                                                                                                                                                                           0; Indels
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                                                                                                 Sequence 12 BP; 3 A; 0 C; 8 G; 1 T; 0 U; 0 Other;
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was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                         12.3%; Score 9; DB 1
100.0%; Pred. No. 1.4
ive 0; Mismatches
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Matches 9; Conservative
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Best Local Similarity
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central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                         Homo sapiens.
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                                                                                                                                                                                                                                                                                                                                                                                                                                               This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99889, ABF00010-ABF99889, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                               SNP, single nucleotide polymorphism, human; diagnosis, PNA, cancer, CNS, peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss;
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                                                                                                         Oligonucleotide primer SEQ ID NO 367643 for detecting SNP TSC0004601.
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                                      ABI67670 standard; DNA; 12 BP.
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                                                                                    (first entry)
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Best Local Similarity 100.
Matches 9; Conservative
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               RESULT 2670
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABIC0010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                                                                                                                                                                                                                                                                                       Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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06-APR-2001; 2001WO-IB000713.
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                                                                                                                                 ABI18422 standard; DNA; 12
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5. Conservative
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         WPI; 2001-657177/75
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                                                                                                                                       ABI18422;
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and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                        12.3%; Score 9; DB 1; Length 12; 100.0%; Pred. No. 1.4e+03; ive 0; Mismatches 0; Indels
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les 9; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                            Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                                                                                                                     Claim 1; SEQ ID NO 363780; 29pp + Sequence Listing; German.
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                            Berlin K;
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                            Piepenbrock C,
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically prereated genomic DNA. The oligomuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABE99989, ABF00010-ABE99989, ABH00010-ABE99989 and ABI00010-ABE92073 represent the oligomers described in the invention. NOTE: The sequence was obtained in electronic formmat from WIPO at the printed specification, but the wipo.int/pub/published_pct_sequences Sequence 12 BP; 3 A; 3 C; 0 G; 6 T; 0 U; 0 Other;

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP)

Claim 1; SEQ ID NO 318395; 29pp + Sequence Listing; German.

oet or oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.

WPI; 2001-657177/75.

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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, contral nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 trepresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                    SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; 88; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                   Oligonucleotide primer SEQ ID NO 273338 for detecting SNP TSC0003145.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 1; SEQ ID NO 273338; 29pp + Sequence Listing; German.
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                                                                                                                                                                                                                                                                                                                     07-APR-2000; 2000DE-01019173.
                                   (first entry)
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 ABH73353;
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Matches
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                                                                                                                                                                                                                                                                                                                                  SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                    Oligonucleotide primer SEQ ID NO 296172 for detecting SNP TSC0016943.
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12.3%; Score 9; DB 1; Length 12; 100.0%; Pred. No. 1.4e+03; ive 0; Mismatches 0; Indels
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                                                           12.3%; Score 9; DB 1; Length 12; ilarity 100.0%; Pred. No. 1.48+03; Conservative 0; Mismatches 0: Indela
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acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomucleotides are used for diagnosis and/or prognosis of cancer and range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC09989, ABC0010-ABE9989, ABC0010-ABE9989, ABC0010-ABE9989, ABC0010-ABE9989 and ABI0010-ABE82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                                                                                                                                                                     Claim 1; SEQ ID NO 301138; 29pp + Sequence Listing; German.
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Best Local Similarity
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nuclectide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The coligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The coligomers are also used for detecting cell type differentiation. ABC0010-ABC99999, ABF00010-ABF99999, ABH00010-ABH99999 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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diagnosis and cell typing, is
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                designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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Piepenbrock C,

olek A,

WPI; 2001-657177/75

(EPIG-) EPIGENOMICS AG

primers or peptide nucleic

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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                 Oligonucleotide primer SEQ ID NO 342165 for detecting SNP TSC0042413.
                                                                                           ABI42192 standard; DNA; 12 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                       SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pot_sequences
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12.3%; Score 9; DB 1; Length 12;
Best Local Similarity 100.0%; Pred. No. 1.46+03;
Matches 9; Conservative 0; Mismatches 0; Indels
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PMA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF99989, ABH00010-ABF99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but twipo.int/pub/published_pct_sequences
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, contral nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                           SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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nes 9; Conservative
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF99989, ABH00010-ABF99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic formmat from WIPO at
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07-APR-2000; 2000DE-01019173.
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                                                             Piepenbrock C,
                              (EPIG-) EPIGENOMICS AG
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Claim 1; SEQ ID NO 370307; 29pp + Sequence Listing; German.

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                       This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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Sequence 12 BP; 4 A; 0 C; 1 G; 7 T; 0 U; 0 Other;
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AB16
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RESULT 2687

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF99999, ABH00010-ABH99999 and ABI00010-AB182073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic formmat from WPPO at fib. wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Oligonucleotide primer SEQ ID NO 273147 for detecting SNP TSC0003063.
                                                                                                                                                                                                                                                                                                                                             oligonucleotides, useful for diagnosis and cell typing, ied to detect single-nucleotide polymorphisms and cytosine
                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; SEQ ID NO 294341; 29pp + Sequence Listing; German.
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                                                                                                                                                                                                                                                            Berlin K;
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                                                                                                                                                                                                                                                          Piepenbrock C,
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                                                                                                                                                                                                                                                                                                                                                                                           methylation status.
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  Homo sapiens.
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                                                                                                                                                                                           SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                    Oligonucleotide primer SEQ ID NO 366269 for detecting SNP TSC0055636
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                    ABI66296 standard; DNA; 12
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Matches 9; Conservative
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genemic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99899, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Oligonucleotide primer SEQ ID NO 282790 for detecting SNP TSC0010992.
                                                      Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                     Claim 1; SEQ ID NO 273147; 29pp + Sequence Listing; German.
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range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ASC0010-ABC09989, ABC0010-ABE9989, ABH0010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence was obtained in electronic format from WIPO at the printed specification, but the wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                       Seguence 12 BP; 6 A; 0 C; 3 G; 3 T; 0 U; 0 Other;
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nuclectide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic discorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABC00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but twipo.int/pub/published_pct_sequences
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                                                                                     SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                          Oligonucleotide primer SEQ ID NO 369605 for detecting SNP TSC0057741.
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designed to detect single-nucleotide polymorphisms and cytosine
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                                                                                                                                                                                                                                                                                                                                                                                                     (EPIG-) EPIGENOMICS AG
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Matches
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                                                                                                                                                                                                                                                                                                                                                                              SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; 88; central nervous system; gastrointestinal; respiratory; immune; metabolic.
    Gaps
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                                                                                                                                                                                                                                                                                                                                     Oligonucleotide primer SEQ ID NO 350967 for detecting SNP TSC0047021.
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    Indels
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    0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Berlin K;
                                                                                                                                                                                                    ABIS0994 standard; DNA; 12 BP.
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                                                 958 CGCTACCAA 966
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tes 9; Conserv
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Gaps

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ABI69632

ABI69632 ID ABI XX AC ABI

RESULT 2693

Matches

8 g

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methylation status.
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                                                                                                                                                                                                                                 This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                   Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                                                                                                                        Claim 1; SEQ ID NO 378587; 29pp + Sequence Listing; German.
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                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 12 BP; 3 A; 5 C; 0 G; 4 T; 0 U; 0 Other;
             06-APR-2001; 2001WO-IB000713.
                                        07-APR-2000; 2000DE-01019173
                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 100.
Matches 9; Conservative
                                                                                             Olek A, Piepenbrock C,
                                                                    (EPIG-) EPIGENOMICS AG
                                                                                                                         WPI; 2001-657177/75.
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Oligonucleotide primer SEQ ID NO 366588 for detecting SNP TSC0055854. ABI66615 standard; DNA; 12 BP. (first entry) 22-FEB-2002 ABI66615; 2695

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Homo sapiens

WO200177384-A2.

18-OCT-2001

06-APR-2001; 2001WO-IB000713

07-APR-2000; 2000DE-01019173.

(EPIG-) EPIGENOMICS AG

Berlin K; Olek A, Piepenbrock C,

WPI; 2001-657177/75.

Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine

Claim 1; SEQ ID NO 366588; 29pp + Sequence Listing; German.

This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at

Sequence 12 BP; 5 A; 3 C; 1 G; 3 T; 0 U; 0 Other;

0; Gaps 12.3%; Score 9; DB 1; Length 12; 100.0%; Pred. No. 1.4e+03; ative 0; Mismatches 0; Indels Query Match
Best Local Similarity 100.
Matches 9, Conservative

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947 GITTAATGT 955 σı ABI17925 standard; DNA; 12 BP. (first entry) 22-FEB-2002 ABI17925;

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Gaps

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Oligonucleotide primer SEQ ID NO 317898 for detecting SNP TSC0028334.

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Homo sapiens.

WO200177384-A2.

18-OCT-2001

06-APR-2001; 2001WO-IB000713.

07-APR-2000; 2000DE-01019173.

(EPIG-) EPIGENOMICS AG.

Berlin K; Olek A, Piepenbrock C,

WPI; 2001-657177/75.

set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.

Claim 1; SEQ ID NO 317898; 29pp + Sequence Listing; German.

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABE03989, ABE00010-ABE9989, ABE00010-ABE9989 and ABI00010-ABE82073 represent the oligomers described in the invention. NOTE: The sequence

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RESULT 26
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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                                                                                                                                                    12.3%; Score 9; DB 1; Length 12; 100.0%; Pred. No. 1.4e+03; tive 0; Mismatches 0; Indels
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Best Local Similarity 100.
Matches 9; Conservative
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABF3073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but typ.wipo.int/pub/published_pot_sequences
                                                                                                                                                                                                                       SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                 Oligonucleotide primer SEQ ID NO 277861 for detecting SNP TSC0005098
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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ABH77868 standard; DNA; 12
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Best Local Similarity 100.
Matches 9; Conservative
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS;

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Gaps

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0; Indels

9; Conservative

Matches

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(EPIG-) EPIGENOMICS AG
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                                                                                                                                                                                                                                                                            This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, contral nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF9989, ABF00010-ABF9989, and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
 nucleic acid, cytosine methylation; cardiovascular; primer; ss; nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                         Claim 1; SEQ ID NO 306620; 29pp + Sequence Listing; German.
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Matches 9; Conservative
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                                                                                                                                              (EPIG-) EPIGENOMICS AG
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                                  Homo sapiens.
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nuclectide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI32073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                    set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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12.3%; Score 9; DB 1; Length 12;
Best Local Similarity 100.0%; Pred. No. 1.4e+03;
Matches 9; Conservative 0; Mismatches 0; Indels
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Olek A, Piepenbrock C,
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acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC09989, ABF00010-ABE09989, ABF00010-ABE09989, ABF00010-ABE09989, ABF00010-ABE09989, DABF00010-ABE09989, ABF00010-ABE09989, ABF00010-ABE09998, ABF00010010-ABE09998, ABF00010-ABE09998, ABF00010-ABE09998, ABF000010-ABE09998, ABF00010-ABE09998, ABF00010-ABE09998, ABF00010-ABE09998, 
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                                                                                                                                                                                                                                                                                                                                 SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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designed to detect single-nucleotide polymorphisms and cytosine
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 DB 1; Length 12;
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iive 0; Mismatches
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                Best Local Similarity 100.
Matches 9; Conservative
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1563/c ABH91563 standard; DNA; 12 BP.

RESULT 2704 ABH91563/c ID ABH9156:

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF9989, ABF00010-ABF9989, ABF00010-ABF9989, ABF00010-ABF9989, and ABI00010-ABF8003 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at

Sequence 12 BP; 6 A; 4 C; 0 G; 2 T; 0 U; 0 Other;

Claim 1; SEQ ID NO 288951; 29pp + Sequence Listing; German.

ABH91563;

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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                  Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                    SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                 Oligonucleotide primer SEQ ID NO 291556 for detecting SNP TSC0014831.
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ftp.wipo.int/pub/published_pct_sequences
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
Oligonucleotide primer SEQ ID NO 344908 for detecting SNP TSC0043756
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Homo sapiens

ABI42422;

RESULT 2705

Query Match

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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretraeted genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99899, ABF00010-ABF99989 and ABF00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
          Set of oligonucleotides, useful for diagnosis and cell typing, : designed to detect single-nucleotide polymorphisms and cytosine
                                                                   Claim 1; SEQ ID NO 344908; 29pp + Sequence Listing; German.
                                                                                                                                                                                                                                                                                                         Query Match 12.3%; Score 9; DB 1; Length 12; Best Local Similarity 100.0%; Pred. No. 1.4e+03; Matches 9; Conservative 0; Mismatches 0; Indels
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                                      methylation status.
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ABI49477 standard; DNA; 12 BP.
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic. Oligonucleotide primer SEQ ID NO 349450 for detecting SNP TSC0046149,

Berlin K;

WPI; 2001-657177/75.

bet of oligonuclectides, useful for diagnosis and cell typing, addesigned to detect single-nuclectide polymorphisms and cytosine methylation status.

Claim 1; SEQ ID NO 349450; 29pp + Sequence Listing; German.

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The

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oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF0010-ABF9989, ABF0010-ABF9989, ABF0010-ABF9989, ABF0010-ABF9989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at flow the printed in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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ative 0; Mismatches
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nes 9; Conservative
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Oligonucleotide primer SEQ ID NO 298649 for detecting SNP TSC0018216.
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                                                                                                                                                                                                                                                                                                                                                                              SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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12.3%; Score 9; DB 1; Length 12;
Best Local Similarity 100.0%; Pred. No. 1.4e+03;
Matches 9; Conservative 0; Mismatches 0; Indels
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          943 ATTGGTTTA 951
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and oytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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SND; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Berlin

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form par to f the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                                                   onucleotides, useful for diagnosis and cell typing, i detect single-nucleotide polymorphisms and cytosine
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                                                                                                                                                                                                                     Claim 1; SEQ ID NO 301399; 29pp + Sequence Listing; German.
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Local Similarity 100.0%; Pred. No. 1.4e+03;
les 9; Conservative 0; Mismatches 0; Indels
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                 07-APR-2000; 2000DE-01019173
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                                                                                                                                                   of oligonucleotides,
                                                                                 Piepenbrock C,
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                                                                                                                   WPI; 2001-657177/75
                                                                                                                                                                   designed to detect methylation status.
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Berlin K;

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PMA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABE99989, ABF00010-ABE99989, ABH00010-ABH99989 and ABI00010-ABE82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at
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Claim 1; SEQ ID NO 327213; 29pp + Sequence Listing; German.
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                Oligonucleotide primer SEQ ID NO 329815 for detecting SNP TSC0035174.
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                          ABI29842 standard; DNA; 12
                                                                                     (first entry)
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                                                                                                                                                                                                                                                                                                                                        SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                      Query Match 12.3%; Score 9; DB 1; Length 12; Best Local Similarity 100.0%; Pred. No. 1.46+03; Matches 9; Conservative 0; Mismatches 0; Indels
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                            Sequence 12 BP; 3 A; 0 C; 8 G; 1 T; 0 U; 0 Other;
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Best Local Similarity 100.0%; Pred. No. 1.4
Matches 9; Conservative 0; Mismatches
ftp.wipo.int/pub/published_pct_sequences
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Berlin K;

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABF99989, ABH00010-ABF99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic formmat from WIPO at
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Best Local Similarity 100.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPD at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Oligonuclectide primer SEQ ID NO 337831 for detecting SNP TSC0040087.
                                                                Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                   Claim 1; SEQ ID NO 287711; 29pp + Sequence Listing; German.
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nes 9; Conservative
 Piepenbrock C,
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methylation status.
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                                                                                                                                                                                                                                                                                                                                                                            This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nuclectide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                               Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                   Homo sapiens.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The

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oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99889, ABF00010-ABF99889, ABH00010-ABH99889 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at fig. wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                            12.3%; Score 9; DB 1; Length 12; 100.0%; Pred. No. 1.46+03; tive 0; Mismatches 0; Indels
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Best Local Similarity 100.
Matches 9; Conservative
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                                                                                                                                                                                                                                                                                                    SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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designed to detect single-nucleotide polymorphisms and cytosine
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                Indels
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tive 0; Mismatches 0;
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Matches 9; Conservative
                9; Conservative
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                                                930 ATCCCTCCT 938
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Best Local Similarity
Matches 9; Conserv
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Berlin K;

Piepenbrock C,

olek A,

WPI; 2001-657177/75

(EPIG-) EPIGENOMICS AG

06-APR-2001; 2001WO-IB000713. 07-APR-2000; 2000DE-01019173.

18-OCT-2001.

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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                              Oligonucleotide primer SEQ ID NO 355346 for detecting SNP TSC0007599.
                                                                                                                                                                                                                                                                ligonucleotides, useful for diagnosis and cell typing, : to detect single-nucleotide polymorphisms and cytosine
                                                                                                                                                                                                                                                                                                          Claim 1; SEQ ID NO 355346; 29pp + Sequence Listing; German.
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                                                                                                                                                         06-APR-2001; 2001WO-IB000713.
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          (first entry)
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                                                                                             Homo sapiens.
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nuclectide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99899, ABF00010-ABF99899, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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12.3%; Score 9; DB 1; Ler 100.0%; Pred. No. 1.4e+03; trive 0; Mismatches 0; Local Similarity 100. Query Match Matches 6

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ABI61962 standard; DNA; 12 BP ABI61962;

22-FEB-2002 (first entry)

Oligonucleotide primer SEQ ID NO 361935 for detecting SNP TSC0007080

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

Homo sapiens

WO200177384-A2

Set of oligonucleotides, useful for diagnosis and cell typing, is

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF9989, ABH00010-ABH9989 and ABI00010-ABI82073 the present the oligomers described in the invention. NOTE: The sequence data for this patent din ot form part of the printed specification, but was obtained in electronic format from WIPO at
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designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                         Claim 1; SEQ ID NO 293607; 29pp + Sequence Listing; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12.3%; Score 9; DB 1; Length 12; 100.0%; Pred. No. 1.4e+03; tive 0; Mismatches 0; Indels
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Best Local Similarity 100...
9; Conservative
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represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at from to int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                                                                                                  Oligonucleotide primer SEQ ID NO 320746 for detecting SNP TSC0029862.
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ches 0; Indels
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                                                                                         Sequence 12 BP; 6 A; 0 C; 3 G; 3 T; 0 U; 0 Other;
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                                                                                                                         Query Match
12.3%; Score 9; DB 1
Best Local Similarity 100.0%; Pred. No. 1.4
Matches 9; Conservative 0; Mismatches
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Les 9; Conservative
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF9989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                               SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                   Oligonucleotide primer SEQ ID NO 327214 for detecting SNP TSC0033501.
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100.0%; Pred. No. 1.4e+03;
live 0; Mismatches 0; Indels
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                                 ABI27241 standard; DNA; 12 BP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            methylation status.
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF9989, ABH00010-ABH9989 and ABI00010-ABI82073 invepresant the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                  Berlin K;
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Best Local Similarity 100.
Matches 9; Conservative
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ABH84492
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07-APR-2000; 2000DE-01019173

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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABE99989, ABF00010-ABE99989, ABH00010-ABE99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Oligonuclectide primer SEQ ID NO 286156 for detecting SNP TSC0012603.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12.3%; Score 9; DB 1; Length 12; 100.0%; Pred. No. 1.4e+03; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 12 BP; 7 A; 0 C; 3 G; 2 T; 0 U; 0 Other;
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                       This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at the printed specification, but fire wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Oligonucleotide primer SEQ ID NO 336225 for detecting SNP TSC0039257.
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to detect single-nuclectide polymorphisms and cytosine
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                                                                                                                                                                                                                                                                                                                                                  Claim 1; SEQ ID NO 284485; 29pp + Sequence Listing; German.
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                                                                                                        Berlin K;
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Best Local Similarity 100...
3, Conservative
                                                                                                                                                                                                                             Set of oligonucleotides,
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                                                                                                        Piepenbrock C,
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                                                                                                                                                                                                                                                          designed to detect amethylation status.
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acid (DNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABF9989 and ABT00010-ABF9803, represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the, wipo.int/pub/published_pct_sequences
                                                                                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                  Oligonucleotide primer SEQ ID NO 358515 for detecting SNP TSC0051168.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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ABI58542 standard; DNA; 12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABF99989, ABF001010-ABF99989, ABH001010-ABF99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                            Oligonucleotide primer SEQ ID NO 287638 for detecting SNP TSC0013177.
                                                                                           Gaps
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                                         Query Match 12.3%; Score 9; DB 1; Length 12; Best Local Similarity 100.0%; Pred. No. 1.4e+03; Matches 9; Conservative 0; Mismatches 0; Indels
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Sequence 12 BP; 1 A; 4 C; 0 G; 7 T; 0 U; 0 Other;
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Best Local Similarity
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligomucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and merabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                 Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                                                                             Claim 1; SEQ ID NO 317446; 29pp + Sequence Listing; German.
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100.0%; Pred. No. 1.4
:ive 0; Mismatches
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WPI; 2001-657177/75.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for ABH00010-ABH99989 and ABI00010-ABH99989, ABF00010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the was obtained in electronic format from WIPO at
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 1; SEQ ID NO 380583; 29pp + Sequence Listing; German.
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XC ABBIT 7473;
XX Z2-FEB-2002 (first entry)
XX SNP; single nuclectide polymorph,
XW SNP; single nuclectide polymorph,
XW SNP; single nuclectide polymorph,
XW SNP; single nuclectide gastroin mix SN Central nervous system; gastroin mix XX
XX SN WO200177384-A2.
XX PN WO200177384-A2.
XX PF 06-APR-2000; 2000DB-01019173.
XX (EPIG-) EPIGENOMICS AG.
XX X EPIG-)
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945 TGGTTTAAT 953

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central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC09989, ABF00010-ABF9989 and ABI00010-ABF82073 represent the oligomers described in the invention. NoTE: The sequence data for this parent did not form part of the printed specification, but was obtained in electronic format from WIPO at fig. wipo.int/pub/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                                                        SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                                                                      Oligonucleotide primer SEQ ID NO 312504 for detecting SNP TSC0025096.
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designed to detect single-nucleotide polymorphisms and cytosine
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                                                                  ABI39602 standard; DNA; 12 BP.
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22-FEB-2002 (first entry)

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Query Match 12.3%; Score 9; DB 1; Length 12; Best Local Similarity 100.0%; Pred. No. 1.4e+03; Matches 9; Conservative 0; Mismatches 0; Indels

Sequence 12 BP; 5 A; 3 C; 0 G; 4 T; 0 U; 0 Other;

Homo sapiens.

18-OCT-2001

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABE99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 trepresent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from MIPO at
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                                                                                                                                                                                                                                                                                                                                   Claim 1; SEQ ID NO 352259; 29pp + Sequence Listing; German.
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12.3%; Score 9; DB 1; Length 12;
Best Local Similarity 100.0%; Pred. No. 1.4e+03;
Matches 9; Conservative 0; Mismatches 0; Indels
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06-APR-2001; 2001WO-IB000713.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                     SNP; single nuclectide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                         Oligonucleotide primer SEQ ID NO 339575 for detecting SNP TSC0041078.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Berlin K;
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Best Local Similarity
Matches 9; Conserv
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Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.

WO200177384-A2

18-OCT-2001

22-FEB-2002

ABI52286;

8 a Sequence 12 BP; 1 A; 1 C; 0 G; 10 T; 0 U; 0 Other;

was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABE039989, ABE0010-ABE99989 and ABI0010-ABE92073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but
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                                                                                                                                                                                                                              943 ATTGGTTTA 951
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                                                                   This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretracted genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99899, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form mar of the printed specification, but the was obtained in electronic format from WIPO at
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Claim 1; SEQ ID NO 375080; 29pp + Sequence Listing; German.
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Matches 9; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Oligonucleotide primer SEQ ID NO 379542 for detecting SNP TSC000620.
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   DB 1; Length 12; . 1.4e+03;
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Query Match
12.3%; Score 9; DB 1
Best Local Similarity 100.0%; Pred. No. 1.4
Matches 9; Conservative 0; Mismatches
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Matches 9; Conservative
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                                                                                                                                          907 ATTITITIT 915
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central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                           SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                       Oligonucleotide primer SEQ ID NO 380027 for detecting SNP TSC0063601.
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Best Local Similarity 100.
Matches 9; Conservative
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peptide nucleic acid: ~
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically prereated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99999, ABF00010-ABF99999, ABH00010-ABF99999 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at fixed in the published_pct_sequences
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tive 0; Mismatches 0; Indels
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Berlin K;

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genemic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ACC0010 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from MIPO at
                                               designed to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                     Claim 1; SEQ ID NO 318494; 29pp + Sequence Listing; German.
                                                                                                                                                                                                                                                                             Sequence 12 BP; 5 A; 1 C; 3 G; 3 T; 0 U; 0 Other;
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            Piepenbrock C,
                                  WPI; 2001-657177/75
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            olek A,
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Oligonucleotide primer SEQ ID NO 295787 for detecting SNP TSC0016735.
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12.3%; Score 9; DB 1; Length 12; 100.0%; Pred. No. 1.4e+03; rative 0; Mismatches 0; Indels
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                                                                                       904 GTCATTTTC 912
                                                                                                                 12 GTCALTITC
                         Best Local Similarity
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.

06-APR-2001; 2001WO-IB000713

07-APR-2000; 2000DE-01019173

(EPIG-) EPIGENOMICS AG

Berlin K; Olek A, Piepenbrock C,

WPI; 2001-657177/75.

Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine methylation status.

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genemic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, ardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC9989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 the represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at

Sequence 12 BP; 5 A; 0 C; 5 G; 2 T; 0 U; 0 Other;

was obtained in electronic format from Wl ftp.wipo.int/pub/published_pct_sequences

Claim 1; SEQ ID NO 295787; 29pp + Sequence Listing; German.

This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP)

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and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99999, ABF00010-ABF99989, ABF00010-ABF99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence was obtained in electronic format from WIPO at the printed specification, but the wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99899, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                   SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                          Oligonucleotide primer SEQ ID NO 283498 for detecting SNP TSC0011348.
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designed to detect single-nucleotide polymorphisms and cytosine
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DB 1; Length 12;
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ches 0; Indels
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12.3%; Score 9; DB 1
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tive 0; Mismatches
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ABH83505 standard; DNA; 12 BP.
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Berlin K;
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                               (EPIG-) EPIGENOMICS AG
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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nuclectide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99899, ABH00010-ABH99989 and ABI00010-ABI82073 the persent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                             Claim 1; SEQ ID NO 355125; 29pp + Sequence Listing; German.
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ftp.wipo.int/pub/published_pct_sequences
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acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010

WPI; 2001-657177/75

This invention describes novel oligonucleotide primers or peptide nucleic

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-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this parent aid not form part of the printed specification, but was obtained in electronic format from WIPO at ftp.wipo.int/pub/published_pct_sequences
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Best Local Similarity 100.0%; Pred. No. 1.4e+03;
Matches 9; Conservative 0; Mismatches 0; Indels
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretraeted genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99889, ABF00010-ABF99899, ABF00010-ABF99989, ABF00010-ABF99989, and ABI00010-ABF82073 teprement the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                    SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; 8s; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, contral nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABF99899 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo.int/pub/published_pct_sequences
                                             SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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ABH81368 standard; DNA; 12 BP. ABH81368; ABH81368/ XX AC ABH8: XX AC ABH8: XX ABH8: XX ABH8: XX Oliganian SNP; XW PEPUT: XW PETUT: X

22-FEB-2002 (first entry)

SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic. Oligonucleotide primer SEQ ID NO 281361 for detecting SNP TSC0009680.

Homo sapiens

WO200177384-A2

18-OCT-2001

06-APR-2001; 2001WO-IB000713.

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic. Oligonucleotide primer SEQ ID NO 347903 for detecting SNP TSC0045335 06-APR-2001; 2001WO-IB000713. WO200177384-A2 Homo sapiens. 18-OCT-2001.

ВЪ.

ABI47930 standard; DNA; 12

RESULT 2757

ABI47930

22-FEB-2002 (first entry)

ABI47930;

07-APR-2000; 2000DE-01019173.

Berlin K; Olek A, Piepenbrock C, (EPIG-) EPIGENOMICS AG WPI; 2001-657177/75

Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine methylation status.

Claim 1; SEQ ID NO 347903; 29pp + Sequence Listing; German.

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF99989, ABF00010-ABF99989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence was obtained in electronic format from WIPO at the printed specification, but the wipo.int/pub/published_pct_sequences
      This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, contral nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC9989, ABF00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
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                                                                                                                                                                                                                                                                                                                                                               SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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          Sequence 12 BP; 4 A; 0 C; 3 G; 5 T; 0 U; 0 Other;
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                                                                                                                                                                                                                                                                                                            SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                    ABI81772 standard; DNA; 12
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABE99989, ABF00010-ABE99989, ABF00010-ABE99989, ABF00010-ABE99989 and ABI00010-ABI32073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                                                                                                            acid (PNA) oligomers for detecting single nucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for disponsis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABF9989, ABF00010-ABF9989 and ABI00010-ABF32073 represent the oligomers described in the invention. NoTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from MIPO at
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                     Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine methylation status.
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                                                                                                                    Claim 1; SEQ ID NO 321801; 29pp + Sequence Listing; German.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically prereated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99889, ABF00010-ABF99899, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at fib. wipo.int/pub/published_pct_sequences
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range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABC99989, ABF00010-ABF9989, ABF00010-ABF9989 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this parent did not form part of the printed specification, but was obtained in electronic format from WIPO at the printed specification, but the wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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Best Local Similarity 100.0%; Pred. No. 1.4e+03;
Matches 9; Conservative 0; Mismatches 0; Indels
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but ftp.wipo.int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                      SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                       Oligonucleotide primer SEQ ID NO 304815 for detecting SNP TSC0021123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine methylation status.
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, cointral nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99989, ABF00010-ABF99989, ABH00010-ABH99989 and ABI00010-ABI32073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                             SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
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                                                             Oligonucleotide primer SEQ ID NO 338555 for detecting SNP TSC0040547.
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ABI38582 standard; DNA; 12 BP.

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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABE99989, ABF00010-ABE99989, ABH00010-ABE99989 and ABI00010-ABI82273 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at
                                                                                                                                                                             Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status.
                                                                                                                                                                                                                                           Claim 1; SEQ ID NO 314991; 29pp + Sequence Listing; German.
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                                                                                                                Olek A, Piepenbrock C,
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Best Local Similarity 100.0%; Pred. No. 1.4e+03;
Matches 9; Conservative 0; Mismatches 0; Indels
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic. Oligonucleotide primer SEQ ID NO 344948 for detecting SNP TSC0043792. ABI44975 standard; DNA; 12 BP. 22-FEB-2002 (first entry) Homo sapiens

Berlin K; 07-APR-2000; 2000DE-01019173 06-APR-2001; 2001WO-IB000713 Olek A, Piepenbrock C, (EPIG-) EPIGENOMICS AG. WO200177384-A2. 18-OCT-2001.

Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine

WPI; 2001-657177/75.

This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and expeciate methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABCO010-ABC09989, ABF00010-ABF99989, ABH0010-ABH99989 and ABI00010-ABH82073 represent the oligomers described in the invention. NOTE: The sequence ö This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99889, ABF00010-ABF99989, ABH0010-ABF99989 and ABI00110-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format from WIPO at SNP, single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic. Oligonucleotide primer SEQ ID NO 353754 for detecting SNP TSC0048693. 0; Gaps .H Set of oligonucleotides, useful for diagnosis and cell typing, i designed to detect single-nucleotide polymorphisms and cytosine methylation status. Claim 1; SEQ ID NO 353754; 29pp + Sequence Listing; German. Claim 1; SEQ ID NO 344948; 29pp + Sequence Listing; German. 12.3%; Score 9; DB 1; Length 12; 100.0%; Pred. No. 1.4e+03; ative 0; Mismatches 0; Indels Sequence 12 BP; 4 A; 1 C; 5 G; 2 T; 0 U; 0 Other; Berlin K; ABI53781 standard; DNA; 12 BP. 06-APR-2001; 2001WO-IB000713. 07-APR-2000; 2000DE-01019173. 22-FEB-2002 (first entry) Query Match
Best Local Similarity 100.
Matches 9, Conservative Olek A, Piepenbrock C, (EPIG-) EPIGENOMICS AG. 930 ATCCCTCCT 938 10 ATCCCTCCT 2 WPI; 2001-657177/75 WO200177384-A2. Homo sapiens 18-OCT-2001 ABI53781; RESULT 2769 AB15378: ઠ g

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This invention describes novel oligonuclectide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonuclectides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC00010-ABE99989, ABF00010-ABE99989 and ABI00010-ABI82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo int/pub/published_pct_sequences
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           the printed specification, but
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This invention describes novel oligonucleotide primers or peptide nucleic acid (PNA) oligomers for detecting single nucleotide polymorphisms (SNP) and cytosine methylation status in chemically pretreated genomic DNA. The oligonucleotides are used for diagnosis and/or prognosis of cancer and a range of diseases including immune system, gastrointestinal, respiratory, central nervous system, cardiovascular and metabolic disorders. The oligomers are also used for detecting cell type differentiation. ABC0010-ABC99999, ABF00010-ABF99999, ABH00010-ABF99999 and ABI00010-ABF82073 represent the oligomers described in the invention. NOTE: The sequence data for this patent did not form part of the printed specification, but the wipo int/pub/published_pct_sequences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ;
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                                                                                                                                                                  SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS; peptide nucleic acid; cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                     Oligonucleotide primer SEQ ID NO 366019 for detecting SNP TSC0055490.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Oligonucleotide primer SEQ ID NO 381822 for detecting SNP TSC0064564.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Set of oligonucleotides, useful for diagnosis and cell typing, idesigned to detect single-nucleotide polymorphisms and cytosine
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 1; SEQ ID NO 366019; 29pp + Sequence Listing; German.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  Berlin K;
                                ABI66046 standard; DNA; 12 BP
                                                                                                                                                                                                                                                                                                                                              06-APR-2001; 2001WO-IB000713.
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                                                                                                                                                                                                                                                                                                                                                                               07-APR-2000; 2000DE-01019173
                                                                                                  (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  Piepenbrock C,
                                                                                                                                                                                                                                                                                                                                                                                                                 (EPIG-) EPIGENOMICS AG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      949 TTAATGTAT 957
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      methylation status.
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                                                                                                                                                                                                                                                                        WO200177384-A2
                                                                                                                                                                                                                                           Homo sapiens.
                                                                                                    22-FEB-2002
                                                                                                                                                                                                                                                                                                             18-OCT-2001.
                                                                 AB166046;
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RESULT 2771
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ABI81849
                  ABI66046
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SNP; single nucleotide polymorphism; human; diagnosis; PNA; cancer; CNS;

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0; Gaps

0; Indels

12.3%; Score 9; DB 1; Length 12; 100.0%; Pred. No. 1.4e+03; cive 0; Mismatches 0; Indels

Query Match 12.3 Best Local Similarity 100. Matches 9; Conservative

936 CCTCTTCAT 944 cererrear 11

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peptide nucleic acid, cytosine methylation; cardiovascular; primer; ss; central nervous system; gastrointestinal; respiratory; immune; metabolic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                             Set of oligonucleotides, useful for diagnosis and cell typing, is designed to detect single-nucleotide polymorphisms and cytosine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          JA11 linker DNA used to illustrate the method of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Protein-protein interaction; detection; cancer; linker; ss.
                                                                                                                                                                                                                                                                           Claim 1; SEQ ID NO 381822; 29pp + Sequence Listing; German.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 1; Length 12; . 1.4e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     12.3%; Score 9; DB 1
100.0%; Pred. No. 1.4
:ive 0; Mismatches
                                                                                                                                                                                 Berlin K;
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                                                                                                           06-APR-2001; 2001WO-IB000713.
                                                                                                                                  07-APR-2000; 2000DE-01019173
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Best Local Similarity luv...
3; Conservative
                                                                                                                                                                               Olek A, Piepenbrock C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     951
                                                                                                                                                          (EPIG-) EPIGENOMICS AG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4 ATTGGTTTA 12
                                                                                                                                                                                                      WPI; 2001-657177/75.
                                                                                                                                                                                                                                                       methylation status.
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                                                             WO200177384-A2.
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                                      Homo sapiens
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                                                                                  18-OCT-2001.
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96US-00663824 97US-00874825

14-JUN-1996; 13-JUN-1997;

(CURA-) CURAGEN CORP

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The present invention relates to novel methods for detecting protein to protein interactions amongst two populations of proteins, each having a complexity of at least 100. The method involves using new genetic methods in which encoded proteins are fused to either the DNA-binding domain of a transcriptional activator or the activation domain of a transcriptional activator or the activation domain of a transcriptional activator. The methods are useful to detect interacting proteins and to identify protein interactions specific for a particular species, tissue, stage of development or disease state, e.g. by comparing protein-protein interactions between populations from conditions or precancerous cells with those from non-cancerous cells. They are also useful to identify inhibitors interfering with protein-protein interactions e.g. populations from protein protein interactions e.g. populations from protein protei
                                                                                                                         Detection of protein to protein interactions amongst two protein populations useful e.g. to identify interactions specific for particular tissues or diseases and to identify inhibitors of interactions uses a new genetic method.
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Kalbfleisch TS;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 12 BP; 1 A; 4 C; 3 G; 4 T; 0 U; 0 Other;
   Knight JR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Lustig M;
       Yang M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Human NAT2 mutant C282T probe #2.
                                                                                                                                                                                                                                                                                                                               Example; Col 201; 152pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Waschuetza S, Schnakenberg E,
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30-APR-2002; 2002DE-01019373.
       Nandabalan K, Rothberg JM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADD24746/c
ID ADD24746 standard; DNA; 12
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                                                                           WPI; 2002-654433/70.
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WPI; 1995-358649/46.

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This invention describes a novel diagnostic kit for determining tolerance of pharmaceuticals in humans by determining allelic variability of at least two polymorphisms of a human enzyme involved in phase I and/or II of the detoxification mechanism in a blood, tissue or other human sample, where tolerance is determined from presence or absence of alleles. The kit comprises two pairs of oligonucleotide primers, in which each pair amplifies, by PCR, part of a gene for a human detoxification mechanism. CC associated enzyme. The kit may also contain two further pairs of oligonucleotides, serving as probes for detection of amplified DNA segments, especially where the probes are complementary to a single strand of one allele of the target gene. The probes are labelled with segments, especially where the probes are complementary to a single strand of one allele of the target gene. The probes are labelled with segments (LC:Red040 or LC-Red056 for 5'-labelling) which generate a different signal in the hybridized and non 'hybridized condition. The enzymes detected include NAI2, CYP2D6, CYP1A2, CYP2D4, mEH, TPMT, MTHER, paraoxonase, CYP2O, CYP2C19, CYP2DE, CYP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 12 BP; 3 A; 1 C; 5 G; 3 T; 0 U; 0 Other;
                                       Disclosure; Page 86; 156pp; German.
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0; Gaps
12.3%; Score 9; DB 1; Length 12; 100.0%; Pred. No. 1.4e+03; ative 0; Mismatches 0; Indels
                                                                                              Sense strand of sequencing probe B.
                                                             AAT04326 standard; DNA; 13 BP.
                                                                                   20-MAY-1996 (first entry)
           9; Conservative
                       929 TATCCCTCC 937
                                 ٣
                                11 TATCCTCC
      Best Local Similarity
                                                                        AAT04326;
 Query Match
                                                 RESULT 2775
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Polymerase chain reaction; PCR; primer; amplify; pUC19; T4 DNA ligase; streptavidin-coated magnetic beads; type II restriction endonuclease; sequence diagnosis; genetic mapping; forensic analysis; probe; ds. recognition site" /note= "4 bp 5' overhang" Location/Qualifiers
7. 12
/*tag= a /note= "Ear I recognic complement (13) misc_feature WO9527080-A2 misc_feature

Synthetic

(LYNX-) LYNX THERAPEUTICS INC

Brenner S;

94US-00222300. 94US-00280441. 95WO-US003678

25-JUL-1994;

04-APR-1994;

24-MAR-1995;

12-OCT-1995

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This sequence represents the sense strand of a sequencing probe of the invention. The antisense strand of this probe is AAT04327. The probe is used to sequence a 368 bp fragment of pucls that was amplified by the probe is the probe is a strached to sequence as 86 bp fragment of pucls that was amplified by the probes are attached to streptavidin-coated magnetic beads, and digested to produce a 5' overhang. The immobilised sequence is then incubated with this set of ligated to the immobilised sequence using T4 DNA ligase. The probes are ligated to the immobilised sequence using T4 DNA ligase. The probes are ligated to the immobilised of the taxget DNA can be determined. This cleavage shortens the immobilised DNA sequence by one nucleotide. By cycling this reaction, the sequence of the taxget DNA can be determined. This method can be used to sequence DNA at a predetermined genetic locus (that has several possible allelic forms) to determine the zygosity of the individual. It can alles be used in sequence diagnosis, genetic mapping or identification, forensic analysis and research in molecular overlapping bands in a gel. Also, there is no need to generate DNA can be template sequence. The process is readily automated and forest and the process is readily automated and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PCR primer; chromosomal abnormality; abnormality detection; leukaemia; lymphoma; carcinoma; adenocarcinoma; sarcoma; glioma; neuroblastoma; medullablastoma; malignant melanoma; malignant neoglastic condition; ss.
                                                              DNA sequencing by repeated ligation of probe and endonuclease cleavage avoids electrophoretic sepn. of similarly sized fragments, partic. for determining zygosity at a particular locus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  sample
using
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12.3%; Score 9; DB 1; Length 13; 100.0%; Pred. No. 1.5e+03; rive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                        Example 5; Page 28; 67pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            real time monitoring is possible
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 12.3
Best Local Similarity 100.
Matches 9; Conservative
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